



STIC Search Report

Biotech-Chem Library

STIC Database Tracking Number: 188266

TO: Nita M Minnifield
Location: rem-3c01/3c18
Art Unit: 1645
Wednesday, May 10, 2006
Case Serial Number: 08/170344

From: Kristine Hensle
Location: Biotech-Chem Library
REM-1B69
Phone: (571)272-4161

Kristine.Hensle@uspto.gov

Search Notes

Examiner Minnifield,

See attached results. This packet is part 1 of 8.

If you have any questions about this search feel free to contact me at any time.

Thank you for using STIC search services!

Kristine Hensle
Librarian
STIC Biotech/Chem Library
(571)272-4161

*Reviewed
5/11/06
MM*

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STIC-Biotech/ChemLib

188266

From: Pak, Michael
Sent: Tuesday, May 02, 2006 1:15 PM
To: STIC-Biotech/ChemLib
Cc: Minnifield, Nita
Subject: FW: sequence search approval 08/170344

Dear STIC,

Please search the multiple sequence search request set forth below.

Thanks,

Mike Pak

-----Original Message-----

From: Minnifield, Nita
Sent: Sunday, April 30, 2006 3:29 PM
To: Pak, Michael
Subject: sequence search approval 08/170344

Michael,
08/170344
Michael,

This case has been ABN since 01/97; however Applicants petitioned to have it revived and of course it was revived. There are 74 short peptide sequences in the claims. I need an interference sequence search done. Please approve so I can get this case moved (this is a 2 month amdt. case).

STIC

Please do an interference sequence search on SEQ ID NO: 1-74 (all aa sequences) of this application.

Please provide a paper copy of all results.

Thanks,
Minnifield,

Searcher: _____
Searcher Phone: _____
Date Searcher Picked up: _____
Date completed: _____
Searcher Prep Time: _____
Online Time: _____

Type of Search
NA# _____ AA# _____
S/L: _____ Oligomer: _____
Encode/Transl: _____
Structure #: _____ Text: _____
Inventor: _____ Litigation: _____

Vendors and cost where applicable
STN: _____
DIALOG: _____
QUESTEL/ORBIT: _____
LEXIS/NEXIS: _____
SEQUENCE SYSTEM: _____
WWW/Internet: _____
Other (Specify): _____

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GenCore version 5.1.7
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OM protein - protein search, using SW model

Run on: May 5, 2006, 06:24:27 ; Search time 26.75 Seconds
(without alignments)
27.816 Million cell updates/sec

Title: US-08-170-344-1

Perfect score: 48
Sequence: 1 AMFDDPQRR 9

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Issued Patents AA:*
1: /cgn2_6/prodata/1/1aa/5-COMB.pep:*
2: /cgn2_6/prodata/1/1aa/6-COMB.pep:*
3: /cgn2_6/prodata/1/1aa/H-COMB.pep:*
4: /cgn2_6/prodata/1/1aa/PCTUS-COMB.pep:*
5: /cgn2_6/prodata/1/1aa/RE-COMB.pep:*
6: /cgn2_6/prodata/1/1aa/backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	48	100.0	9	US-08-159-339A-225	Sequence 225, App
2	48	100.0	23	US-08-363-586-3	Sequence 3, Appl
3	48	100.0	158	US-09-880-523B-2	Sequence 2, Appl
4	48	100.0	162	US-08-316-239B-3	Sequence 3, Appl
5	48	100.0	162	US-08-316-239B-4	Sequence 4, Appl
6	48	100.0	172	US-08-860-165-14	Sequence 14, Appl
7	48	100.0	172	US-09-359-382-14	Sequence 10, Appl
8	48	100.0	182	US-08-117-083-10	Sequence 10, Appl
9	48	100.0	243	US-09-462-993-1	Sequence 1, Appl
10	48	100.0	266	US-08-860-165-10	Sequence 10, Appl
11	48	100.0	266	US-09-359-382-10	Sequence 10, Appl
12	48	100.0	266	US-09-367-309A-1	Sequence 4, Appl
13	48	100.0	273	US-09-485-885-4	Sequence 10, Appl
14	48	100.0	273	US-09-485-885-10	Sequence 10, Appl
15	48	100.0	371	US-09-485-885-6	Sequence 14, Appl
16	48	100.0	390	US-09-485-885-14	Sequence 14, Appl
17	44	91.7	30	US-08-363-586-4	Sequence 4, Appl
18	44	91.7	151	US-09-701-080C-18	Sequence 18, Appl
19	43	89.6	20	US-08-934-915-158	Sequence 158, App
20	39	81.2	15	US-07-909-122-2	Sequence 2, Appl
21	36	75.0	15	US-08-075-541D-52	Sequence 52, Appl
22	35	72.9	1161	US-09-543-681A-6486	Sequence 6486, Ap
23	33	68.8	245	US-09-198-452A-37	Sequence 37, Appl
24	33	68.8	332	US-09-252-991A-32932	Sequence 32932, A
25	33	68.8	588	US-09-438-185A-23	Sequence 23, Appl
26	33	68.8	910	US-09-902-540-16309	Sequence 16309, A
27	32	66.7	325	US-09-702-705-1816	Sequence 1816, Ap

28	32	66.7	325	US-09-736-457-1816	Sequence 1816, Ap
29	32	66.7	325	US-09-671-325-1816	Sequence 1816, Ap
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31	32	66.7	401	US-09-631-663A-2	Sequence 2, Appl
32	32	66.7	529	US-09-248-796A-16357	Sequence 7923, Ap
33	32	66.7	620	US-09-489-039A-7923	Sequence 41556, A
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35	31	64.6	271	US-07-857-224B-46	Sequence 42416, A
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43	31	64.6	545	US-10-237-551-157	Sequence 215, App
44	31	64.6	547	US-10-237-551-215	Sequence 215, App
45	31	64.6	547	US-10-237-551-216	Sequence 9511, Ap
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56	30	62.5	275	US-09-270-767-42414	Sequence 29700, A
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58	30	62.5	326	US-09-710-279-3238	Sequence 4528, Ap
59	30	62.5	333	US-09-134-001C-4528	Sequence 4, Appl
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89	30	62.5	3798	US-09-567-969-6	Sequence 6, Appl
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91	30	62.5	3798	US-09-568-486-6	Sequence 6, Appl
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93	30	62.5	3798	US-09-567-969-6	Sequence 6, Appl
94	30	62.5	3798	US-10-014-717-6	Sequence 6, Appl
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102	30	62.5	7257	2	US-09-568-486-5	Sequence 5, Appli	175	29	60.4	1482	2	US-09-940-316B-21	Sequence 21, Appli
103	30	62.5	7257	2	US-09-568-472-5	Sequence 5, Appli	176	29	60.4	1488	2	US-09-410-551B-17	Sequence 17, Appli
104	30	62.5	7257	2	US-09-567-899-5	Sequence 5, Appli	177	29	60.4	1488	2	US-09-940-316B-17	Sequence 17, Appli
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108	29	60.4	29	1	US-08-318-837-53	Sequence 53, Appli	181	29	60.4	1517	2	US-09-940-316B-19	Sequence 19, Appli
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110	29	60.4	66	2	US-09-270-767-48271	Sequence 48271, A	183	29	60.4	1557	2	US-09-320-878-3	Sequence 3, Appli
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112	29	60.4	94	2	US-09-107-433-4834	Sequence 4834, Ap	185	29	60.4	1562	2	US-09-141-908-4	Sequence 4, Appli
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114	29	60.4	106	1	US-08-318-837-52	Sequence 52, Appli	187	29	60.4	1562	2	US-09-793-708-3	Sequence 3, Appli
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119	29	60.4	216	2	US-09-252-991A-26519	Sequence 26519, A	192	29	60.4	1580	1	US-08-804-227C-11	Sequence 11, Appli
120	29	60.4	218	2	US-09-248-796A-25621	Sequence 25621, A	193	29	60.4	1580	1	US-08-804-198-5	Sequence 5, Appli
121	29	60.4	229	2	US-10-104-047-3781	Sequence 3781, Ap	194	29	60.4	1588	1	US-09-410-551B-29	Sequence 29, Appli
122	29	60.4	235	2	US-09-270-767-42481	Sequence 42481, A	195	29	60.4	1588	2	US-09-940-316B-29	Sequence 29, Appli
123	29	60.4	235	2	US-09-248-796A-24725	Sequence 24725, A	196	29	60.4	1588	2	US-09-410-551B-33	Sequence 33, Appli
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125	29	60.4	255	2	US-09-322-409-58	Sequence 58, Appli	198	29	60.4	1605	2	US-09-252-991A-20395	Sequence 20395, A
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127	29	60.4	256	2	US-09-107-433-4151	Sequence 4151, Ap	200	29	60.4	1721	1	US-08-804-227C-3	Sequence 3, Appli
128	29	60.4	274	2	US-09-322-409-53	Sequence 53, Appli	201	29	60.4	1864	1	US-08-804-227C-3	Sequence 3, Appli
129	29	60.4	274	2	US-09-451-527-53	Sequence 53, Appli	202	29	60.4	1875	2	US-10-042-665A-2	Sequence 2, Appli
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132	29	60.4	298	2	US-09-252-991A-26119	Sequence 26119, A	205	29	60.4	2152	2	US-09-603-207-3	Sequence 3, Appli
133	29	60.4	311	1	US-08-318-837-7	Sequence 7, Appli	206	29	60.4	2392	2	US-09-710-262E-1	Sequence 1, Appli
134	29	60.4	314	2	US-09-902-540-14628	Sequence 14628, A	207	29	60.4	2595	2	US-09-036-987A-2	Sequence 2, Appli
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137	29	60.4	397	2	US-10-104-047-2919	Sequence 2919, Ap	210	29	60.4	3170	1	US-08-439-009A-4	Sequence 4, Appli
138	29	60.4	401	2	US-09-949-016-7956	Sequence 7956, Ap	211	29	60.4	3170	2	US-07-642-734C-5	Sequence 5, Appli
139	29	60.4	402	2	US-09-248-796A-19498	Sequence 19498, A	212	29	60.4	3170	2	US-08-439-009A-5	Sequence 5, Appli
140	29	60.4	412	2	US-09-252-991A-18513	Sequence 18513, A	213	29	60.4	3170	2	US-09-036-987A-4	Sequence 4, Appli
141	29	60.4	426	6	5268463-8	Patent No. 5268463	214	29	60.4	3170	2	US-09-370-700-4	Sequence 4, Appli
142	29	60.4	428	6	5432081-9	Patent No. 5432081	215	29	60.4	3170	2	US-09-603-207-4	Sequence 4, Appli
143	29	60.4	454	2	US-09-904-615-165	Sequence 165, App	216	29	60.4	3201	1	US-09-679-279-15	Sequence 15, Appli
144	29	60.4	454	2	US-10-054-988-165	Sequence 165, App	217	29	60.4	3491	1	US-07-642-734C-2	Sequence 2, Appli
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146	29	60.4	457	1	US-08-882-704A-6	Sequence 6, Appli	219	29	60.4	3519	2	US-09-428-517-4	Sequence 4, Appli
147	29	60.4	457	2	US-09-151-957-6	Sequence 6, Appli	220	29	60.4	3562	2	US-09-679-279-13	Sequence 13, Appli
148	29	60.4	501	2	US-09-248-796A-18735	Sequence 18735, A	221	29	60.4	3562	2	US-09-679-279-14	Sequence 14, Appli
149	29	60.4	508	2	US-09-902-540-13975	Sequence 13975, A	222	29	60.4	3567	1	US-07-642-734C-4	Sequence 4, Appli
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155	29	60.4	772	2	US-09-413-814-92	Sequence 92, Appli	228	29	60.4	3739	2	US-09-105-537-33	Sequence 33, Appli
156	29	60.4	772	2	US-09-252-991A-30121	Sequence 30121, A	229	29	60.4	3739	2	US-09-141-908-3	Sequence 3, Appli
157	29	60.4	785	2	US-09-107-532A-3821	Sequence 3821, Ap	230	29	60.4	3739	2	US-09-657-440-2	Sequence 2, Appli
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159	29	60.4	816	2	US-09-583-110-2733	Sequence 2733, Ap	232	29	60.4	3816	2	US-09-428-517-3	Sequence 3, Appli
160	29	60.4	819	2	US-09-107-433-4725	Sequence 4725, Ap	233	29	60.4	3972	2	US-09-914-286-3	Sequence 3, Appli
161	29	60.4	888	2	US-09-697-022-4	Sequence 4, Appli	234	29	60.4	4551	2	US-09-320-878-1	Sequence 1, Appli
162	29	60.4	888	2	US-09-697-022-5	Sequence 5, Appli	235	29	60.4	4551	2	US-09-141-908-2	Sequence 2, Appli
163	29	60.4	1213	2	US-09-413-814-79	Sequence 79, Appli	236	29	60.4	4551	2	US-09-657-440-1	Sequence 1, Appli
164	29	60.4	1269	2	US-09-252-991A-23150	Sequence 23150, A	237	29	60.4	4551	2	US-09-793-708-1	Sequence 1, Appli
165	29	60.4	1297	2	US-09-328-352-6373	Sequence 6373, Ap	238	29	60.4	4572	2	US-10-042-665A-4	Sequence 4, Appli
166	29	60.4	1300	2	US-09-543-681A-4501	Sequence 4501, Ap	239	29	60.4	4613	2	US-09-105-537-31	Sequence 31, Appli
167	29	60.4	1323	2	US-09-489-039A-13945	Sequence 13945, A	240	29	60.4	4881	2	US-09-914-286-5	Sequence 5, Appli
168	29	60.4	1323	2	US-09-252-991A-17932	Sequence 17932, A	241	29	60.4	4928	2	US-09-036-987A-5	Sequence 5, Appli
169	29	60.4	1346	2	US-09-320-878-4	Sequence 320, Appli	242	29	60.4	4928	2	US-09-370-700-5	Sequence 5, Appli
170	29	60.4	1346	2	US-09-105-537-37	Sequence 37, Appli	243	29	60.4	4928	2	US-09-603-207-5	Sequence 5, Appli
171	29	60.4	1346	2	US-09-141-908-5	Sequence 5, Appli	244	29	60.4	5069	2	US-10-042-665A-5	Sequence 5, Appli
172	29	60.4	1346	2	US-09-657-440-4	Sequence 4, Appli	245	29	60.4	5532	2	US-09-914-286-6	Sequence 6, Appli
173	29	60.4	1346	2	US-09-793-708-4	Sequence 4, Appli	246	29	60.4	5588	2	US-09-036-987A-6	Sequence 6, Appli

247	29	60.4	5588	2	US-09-370-700-6	Sequence 6, Appl1	320	28	58.3	360	2	US-09-905-381A-213	Sequence 213, App
248	29	60.4	5588	2	US-09-603-207-6	Sequence 6, Appl1	321	28	58.3	360	2	US-09-906-618-213	Sequence 213, App
249	29	60.4	6239	2	US-09-914-286-4	Sequence 4, Appl1	322	28	58.3	360	2	US-09-906-646-213	Sequence 213, App
250	29	60.4	6396	2	US-09-410-551B-72	Sequence 72, Appl1	323	28	58.3	360	2	US-09-904-462-213	Sequence 213, App
251	29	60.4	6396	2	US-09-940-316B-72	Sequence 72, Appl1	324	28	58.3	360	2	US-09-902-736A-213	Sequence 213, App
252	29	60.4	11877	2	US-09-105-537-6	Sequence 6, Appl1	325	28	58.3	360	2	US-09-906-722A-213	Sequence 213, App
253	28	58.3	7	2	US-09-884-767A-31	Sequence 31, Appl1	326	28	58.3	361	2	US-09-253-991A-32937	Sequence 17504
254	28	58.3	9	2	US-09-220-081-29	Sequence 29, Appl1	327	28	58.3	362	2	US-09-248-796A-17504	Sequence 18299, A
255	28	58.3	9	2	US-09-677-575-29	Sequence 29, Appl1	328	28	58.3	367	2	US-09-248-796A-18299	Sequence 60125, A
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257	28	58.3	14	2	US-08-466-285-3	Sequence 3, Appl1	330	28	58.3	383	2	US-09-485-885-23	Sequence 5555, Ap
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261	28	58.3	72	2	US-09-270-767-41339	Sequence 41339, A	334	28	58.3	396	2	US-09-378-844-2	Sequence 4, Appl1
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270	28	58.3	126	2	US-09-489-039A-9562	Sequence 9562, Ap	343	28	58.3	430	2	US-09-198-452A-497	Sequence 465, App
271	28	58.3	135	2	US-09-252-991A-26046	Sequence 26046, A	344	28	58.3	430	2	US-09-438-182A-465	Sequence 2, Appl1
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274	28	58.3	147	2	US-09-383-586-35	Sequence 35, Appl1	347	28	58.3	446	2	US-09-976-59A-841	Sequence 7099, Ap
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318	28	58.3	360	2	US-09-909-064-213	Sequence 213, App	391	28	58.3				
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398	58.3	1402	2	US-09-712-363-166	Sequence 166, App	471	27	56.2	304	2	US-09-270-767-33347	Sequence 33347, A
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405	58.3	4472	1	US-08-804-227C-2	Sequence 2, Appli	478	27	56.2	331	2	US-08-560-098A-46	Sequence 46, Appli
406	56.2	38	2	US-09-270-767-61607	Sequence 61607, A	479	27	56.2	333	2	US-09-248-796A-19160	Sequence 19160, A
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408	56.2	41	2	US-09-621-011-142	Sequence 142, App	481	27	56.2	354	2	US-09-786-505-4	Sequence 2, Appli
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444	56.2	217	2	US-09-270-767-34713	Sequence 34713, A	517	27	56.2	404	2	US-08-630-915A-24	Sequence 24, Appli
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547	27	56.2	451	2	US-09-344-889-6	Sequence 6, App1	621	27	56.2	490	2	US-09-949-016-9175	Sequence 9175, App
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741	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	814	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
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747	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	820	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
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749	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	822	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
750	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	823	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
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752	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	825	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
753	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	826	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
754	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	827	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
755	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	828	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
756	27	56.2	1345	2	US-09-902-540-10944	Sequence 10944, A	829	26	54.2	184	2	US-09-568-468-14	Sequence 14, Appl1
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832	26	54.2	215	2	US-09-438-185A-121	Sequence 121, App	905	26	54.2	287	2	US-09-303-518D-915	Sequence 915, App
833	26	54.2	225	1	US-08-462-169B-22	Sequence 22, App1	906	26	54.2	287	2	US-09-303-518D-916	Sequence 916, App
834	26	54.2	225	1	US-08-951-822-25	Sequence 25, App1	907	26	54.2	287	2	US-09-303-518D-917	Sequence 917, App
835	26	54.2	225	2	US-09-103-079-22	Sequence 22, App1	908	26	54.2	287	2	US-09-303-518D-918	Sequence 918, App
836	26	54.2	225	2	US-08-705-245-3	Sequence 25, App1	909	26	54.2	287	2	US-09-303-518D-919	Sequence 919, App
837	26	54.2	225	2	US-09-368-951-25	Sequence 25, App1	910	26	54.2	287	2	US-09-303-518D-920	Sequence 920, App
838	26	54.2	225	2	US-09-425-021-22	Sequence 22, App1	911	26	54.2	287	2	US-09-303-518D-921	Sequence 921, App
839	26	54.2	225	2	US-09-229-947-25	Sequence 25, App1	912	26	54.2	287	2	US-09-303-518D-922	Sequence 922, App
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841	26	54.2	225	2	US-09-564-829-16	Sequence 16, App1	914	26	54.2	288	2	US-09-303-518D-924	Sequence 924, App
842	26	54.2	225	2	US-09-490-714-3	Sequence 3, App1	915	26	54.2	288	2	US-09-303-518D-925	Sequence 925, App
843	26	54.2	225	2	US-08-462-159B-2	Sequence 2, App1	916	26	54.2	288	2	US-09-303-518D-926	Sequence 926, App
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846	26	54.2	225	2	US-09-990-444-497	Sequence 497, App	919	26	54.2	298	2	US-09-488-039A-12846	Sequence 12846, A
847	26	54.2	225	2	US-09-997-333-497	Sequence 497, App	920	26	54.2	306	2	US-09-248-796A-20311	Sequence 20311, A
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849	26	54.2	225	2	US-09-601-040A-28	Sequence 28, App1	922	26	54.2	309	2	US-09-543-681A-4860	Sequence 4860, App
850	26	54.2	227	1	US-09-949-016-6736	Sequence 6736, App	923	26	54.2	311	2	US-09-949-016-8142	Sequence 8142, App
851	26	54.2	228	1	US-08-766-982-11	Sequence 11, App1	924	26	54.2	316	2	US-09-328-352-5882	Sequence 5882, App
852	26	54.2	228	2	US-08-944-483-55	Sequence 55, App1	925	26	54.2	317	2	US-08-859-167-6	Sequence 167-6
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857	26	54.2	234	1	US-08-684-862-3	Sequence 3, App1	930	26	54.2	318	2	US-09-109-273-4	Sequence 273-4
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860	26	54.2	242	2	US-09-802-540-16025	Sequence 16025, A	933	26	54.2	318	2	US-09-276-993-6	Sequence 993-6
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863	26	54.2	252	2	US-09-583-110-3754	Sequence 3754, App	936	26	54.2	318	2	US-09-668-499-2	Sequence 499-2
864	26	54.2	255	2	US-09-252-991A-29130	Sequence 29130, A	937	26	54.2	318	2	US-09-723-450-6	Sequence 450-6
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877	26	54.2	286	2	US-09-711-164-385	Sequence 385, App	950	26	54.2	332	2	US-09-543-681A-4829	Sequence 4829, App
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880	26	54.2	287	2	US-09-303-518D-218	Sequence 218, App	953	26	54.2	345	2	US-09-710-279-2034	Sequence 279-2034
881	26	54.2	287	2	US-09-303-518D-220	Sequence 220, App	954	26	54.2	346	2	US-09-489-039A-7810	Sequence 7810, App
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887	26	54.2	287	2	US-09-303-518D-897	Sequence 897, App	960	26	54.2	352	2	US-09-540-238-2560	Sequence 2560, App
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891	26	54.2	287	2	US-09-303-518D-901	Sequence 901, App	964	26	54.2	356	2	US-09-543-681A-4257	Sequence 4257, App
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893	26	54.2	287	2	US-09-303-518D-903	Sequence 903, App	966	26	54.2	357	2	US-09-540-12478	Sequence 12478, A
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895	26	54.2	287	2	US-09-303-518D-905	Sequence 905, App	968	26	54.2	361	2	US-09-724-797-78	Sequence 797-78
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903	26	54.2	287	2	US-09-303-518D-913	Sequence 913, App	976	26	54.2	377	2	US-10-037-417-66	Sequence 66, App1

977 26 54.2 384 2 US-10-104-047-3188 Sequence 3188, Ap
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981 26 54.2 390 2 US-09-252-991A-22367 Sequence 22367, A
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988 26 54.2 403 2 US-09-248-796A-20519 Sequence 20519, A
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995 26 54.2 427 2 US-08-065-844A-2 Sequence 2, Appl
996 26 54.2 427 2 US-10-002-278-2 Sequence 12, Appl
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998 26 54.2 429 1 US-08-339-152A-33 Sequence 33, Appl
999 26 54.2 435 2 US-09-538-092-538 Sequence 538, Appl
1000 26 54.2 435 2 US-09-487-558B-150 Sequence 150, Appl

ALIGNMENTS

RESULT 1
US-08-159-339A-225
Sequence 225, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Eateban
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
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REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 225:

SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-225

Query Match 100.0%; Score 48; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
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Db 1 AMFODPOER 9

RESULT 2
US-08-363-586-3
Sequence 3, Application US/08363586
Patent No. 5629161
GENERAL INFORMATION:
APPLICANT: Mueller, Martin
APPLICANT: Gissmann, Lutz
TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
TITLE OF INVENTION: Peptides for the Diagnostic Purpose
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,586
FILING DATE: 23-DEC-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/909,296
FILING DATE: 09-JUL-1992
APPLICATION NUMBER: EP 9111720.8
FILING DATE: 13-JUL-1991
ATTORNEY/AGENT INFORMATION:
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REFERENCE/DOCKET NUMBER: 02481-1195-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4400
FILING DATE: 202-408-4400
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-363-586-3

Query Match 100.0%; Score 48; DB 1; Length 23;
Best Local Similarity 100.0%; Pred. No. 0.0084;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
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Db 7 AMFODPOER 15

RESULT 3
US-09-980-523A-2
Sequence 2, Application US/09980523A
Patent No. 6783763
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCINE
APPLICANT: FERRIES, ESTELLE
TITLE OF INVENTION: POLYBIOLOGIC PROTEIN FRAGMENTS OF THE E6 AND E7
TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
FILE REFERENCE: WO/99/01513
CURRENT FILING DATE: 2002-04-29
PRIOR APPLICATION NUMBER: PCT/FR00/01513
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: FR 99/07012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match 100.0%; Score 48; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.064;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||
Db 7 AMFODPOER 15

RESULT 4
US-08-316-239B-3
Sequence 3, Application US/08316239B
Patent No. 5679509
GENERAL INFORMATION:
APPLICANT: Wheeler, Cosette M.
TITLE OF INVENTION: Methods and a Diagnostic Aid for
TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
TITLE OF INVENTION: Cervical Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jagtiani & Associates
STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtiani, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-3

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.064;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||
Db 7 AMFODPOER 15

RESULT 5
US-08-316-239B-4
Sequence 4, Application US/08316239B
Patent No. 5679509
GENERAL INFORMATION:
APPLICANT: Wheeler, Cosette M.
TITLE OF INVENTION: Methods and a Diagnostic Aid for
TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
TITLE OF INVENTION: Cervical Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jagtiani & Associates
STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtiani, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-4

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.064;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||
Db 7 AMFODPOER 15

RESULT 6

```
US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: PRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14

Query Match      100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
   |||||
Db 76 AMFODPOER 84

RESULT 7
US-09-359-382-14
; Sequence 14, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: PRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157/94
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-359-382-14

Query Match      100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.068;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
   |||||
Db 76 AMFODPOER 84

RESULT 8
US-08-117-083-10
```

```
; Sequence 10, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Bourneill, Michael E.
; APPLICANT: Ingils, Stephen C.
; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; TITLE OF INVENTION: Papilloma Virus Proteins
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESS: Walter H. Dreyer
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dreyer, Walter H.
; REGISTRATION NUMBER: 24,190
; REFERENCE/DOCKET NUMBER: A-58783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 182 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..182
; OTHER INFORMATION: /note="Xaa refers to stop codon in
; OTHER INFORMATION: the open reading frame."
US-08-117-083-10

Query Match      100.0%; Score 48; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.072;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
   |||||
Db 8 AMFODPOER 16

RESULT 9
US-09-462-993-1
; Sequence 1, Application US/09462993
; Patent No. 6884786
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BILLOU, Jean-Marc
; APPLICANT: BIZOUANE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 01753-122
; CURRENT APPLICATION NUMBER: US/09/462,993
; CURRENT FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
```

NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TWf.
US-09-462-993-1

Query Match 100.0%; Score 48; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.098;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 35 AMFODPOER 43

RESULT 10
US-08-860-165-10
Sequence 10, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Scirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: PRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P0157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10

Query Match 100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 7 AMFODPOER 15

RESULT 11
US-09-359-382-10
Sequence 10, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Scirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: PRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868

EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P0157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 7 AMFODPOER 15

RESULT 12
US-09-367-309A-1
Sequence 1, Application US/09367309A
Patent No. 6428807
GENERAL INFORMATION:
APPLICANT: MACCARLAN, RODERICK I.
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 7 AMFODPOER 15

RESULT 13
US-09-485-885-4
Sequence 4, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Christine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP96/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4

LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 48; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
Db 113 AMFODPOER 121

RESULT 14
US-09-485-885-10
Sequence 10, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 48; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
Db 132 AMFODPOER 140

RESULT 15
US-09-485-885-6
Sequence 6, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien

US-09-485-885-6

Query Match 100.0%; Score 48; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
Db 113 AMFODPOER 121

RESULT 16
US-09-485-885-14
Sequence 14, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 48; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.16;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
Db 132 AMFODPOER 140

RESULT 17
US-08-363-586-4
Sequence 4, Application US/08363586
Patent No. 5629161
GENERAL INFORMATION:
APPLICANT: Mueller, Martin
APPLICANT: Gissmann, Iutz
TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
TITLE OF INVENTION: Peptides for the Diagnostic Purpose
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,586
FILING DATE: 23-DEC-1994
CLASSIFICATION: 435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/909,296
FILING DATE: 09-JUL-1992
APPLICATION NUMBER: EP 9111720.8
FILING DATE: 13-JUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Wadler, Linda A.
REGISTRATION NUMBER: 33,218
REFERENCE/DOCKET NUMBER: 02481-1195-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-363-586-4

Query Match 91.7%; Score 44; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.066;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
|||||
DB 1 MFQDPOER 8

RESULT 18
US-09-701-080C-18
Sequence 18, Application US/09701080C
Patent No. 6864054
GENERAL INFORMATION:

APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REFERENCE: N73477C GCW
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: US/09/701,080C
PRIOR FILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: GB 9900157.0
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 18
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match 91.7%; Score 44; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.36;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
|||||
DB 1 MFQDPOER 8

RESULT 19
US-08-934-915-158
Sequence 158, Application US/08934915
Patent No. 5932412
GENERAL INFORMATION:
APPLICANT: DILLNER, JOAKIM
APPLICANT: CHENG, HWEI-MING
TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
TITLE OF INVENTION: USEFUL IN IMMUNOSSAY FOR

TITLE OF INVENTION: DIAGNOSTIC PURPOSES
NUMBER OF SEQUENCES: 193
CORRESPONDENCE ADDRESS:

ADDRESSEE: MASON & ASSOCIATES, P.A.
STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
CITY: CLEARWATER
STATE: FLORIDA
COUNTRY: U.S.A.

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 3.0
SOFTWARE: Microsoft Word 6.0
CURRENT APPLICATION DATA:

FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 07/949,836
FILING DATE:

ATTORNEY/AGENT INFORMATION:

NAME: LOUISE A. Fouch

REGISTRATION NUMBER: 37,133

REFERENCE/DOCKET NUMBER: 1946.6

TELECOMMUNICATION INFORMATION:

TELEPHONE: 813-538-3800

TELEFAX: 813-538-3820

TELEX:

INFORMATION FOR SEQ ID NO: 158:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: peptide
US-08-934-915-158

Query Match 89.6%; Score 43; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 0.068;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFQDPOER 9
|||||
DB 6 AMFQDPOER 14

RESULT 20

US-07-909-122-2
Sequence 2, Application US/07909122

Patent No. 5415995
GENERAL INFORMATION:

APPLICANT: SCHOOLNIK, GARY K.

APPLICANT: PALEFSKY, JOEL M.

TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA

TITLE OF INVENTION: VIRUS

NUMBER OF SEQUENCES: 17

CORRESPONDENCE ADDRESS:

ADDRESSEE: MORRISON & FOERSTER

STREET: 755 Page Mill Road

CITY: Palo Alto

STATE: California

COUNTRY: USA

ZIP: 94304-1018

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/909,122

FILING DATE: 19920706

CLASSIFICATION: 530

ATTORNEY/AGENT INFORMATION:

NAME: BENZ, WILLIAM H.

REGISTRATION NUMBER: 25,952
REFERENCE/DOCKET NUMBER: 28600-20105.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
US-07-909-122-2

Query Match 81.2%; Score 39; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.3;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
Db 1 FODPOER 7

RESULT 21
US-08-075-541D-52
Sequence 52, Application US/08075541D
Patent No. 6183745

GENERAL INFORMATION:
APPLICANT: TINDLE, ROBERT
APPLICANT: FERNANDO, GERMAIN
APPLICANT: FRAZER, IAN
TITLE OF INVENTION: SUBUNIT PAPILLOMA VIRUS VACCINE AND
TITLE OF INVENTION: PEPTIDES FOR USE THEREIN
NUMBER OF SEQUENCES: 56
CORRESPONDENCE ADDRESS:
ADDRESSEE: PANITCH SCHWARZE JACOBS & NADEL, P.C.
STREET: 1601 MARKET STREET, 36TH FLOOR
CITY: PHILADELPHIA
STATE: PENNSYLVANIA
COUNTRY: USA
ZIP: 19103-2398

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/075,541D
FILING DATE: 10-JUN-1993
CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU pk 3876
FILING DATE: 12-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: pct/au91/00575
FILING DATE: 12-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: NADEL, ALAN S

REGISTRATION NUMBER: 27,363
REFERENCE/DOCKET NUMBER: 8795-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-567-2020
TELEFAX: 215-567-2991
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear

MOLECULE TYPE: peptide
US-08-075-541D-52

Query Match 75.0%; Score 36; DB 2; Length 15;

Best Local Similarity 85.7%; Pred. No. 1.2;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
Db 1 FODPOER 7

RESULT 22
US-09-543-681A-6486
Sequence 6486, Application US/09543681A
Patent No. 6605709

GENERAL INFORMATION:
APPLICANT: GARY BRETON
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543,681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 6486
LENGTH: 1161
TYPE: PRT

ORGANISM: Proteus mirabilis
US-09-543-681A-6486

Query Match 72.9%; Score 35; DB 2; Length 1161;
Best Local Similarity 75.0%; Pred. No. 1.76+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 308 LFODPOER 315

RESULT 23
US-09-198-452A-37
Sequence 37, Application US/09198452A
Patent No. 6559294

GENERAL INFORMATION:
APPLICANT: Griffaile, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragments
TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, prevention
FILE REFERENCE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NO 37
LENGTH: 245
TYPE: PRT

ORGANISM: Chlamydia pneumoniae
US-09-198-452A-37

Query Match 68.8%; Score 33; DB 2; Length 245;
Best Local Similarity 100.0%; Pred. No. 82;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 ODDPOER 9
Db 61 ODDPOER 66

RESULT 24
US-09-252-991A-32932
Sequence 32932, Application US/09252991A
Patent No. 6551795

GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS


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; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252.991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 32932
; LENGTH: 352
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
; US-09-252-991A-32932

Query Match
Best Local Similarity 68.8%; Score 33; DB 2; Length 352;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 8
Db 163 ALFRDPER 170

RESULT 25
US-09-438-185A-23
; Sequence 23, Application US/09438185A
; Patent No. 6822071
; GENERAL INFORMATION:
; APPLICANT: Stephens, Richard
; APPLICANT: Mitchell, Wayne
; APPLICANT: Kalman, Sue
; APPLICANT: Davis, Ronald
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
; FILE REFERENCE: 018941-000411US
; CURRENT APPLICATION NUMBER: US/09/438,185A
; CURRENT FILING DATE: 2002-03-13
; PRIOR APPLICATION NUMBER: US 60/108,279
; PRIOR FILING DATE: 1998-11-12
; PRIOR APPLICATION NUMBER: US 60/128,606
; PRIOR FILING DATE: 1999-04-08
; NUMBER OF SEQ ID NOS: 1074
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 23
; LENGTH: 588
; TYPE: PRT
; ORGANISM: Chlamydia pneumoniae
; FEATURE:
; OTHER INFORMATION: Cpn0021
; US-09-438-185A-23

Query Match
Best Local Similarity 68.8%; Score 33; DB 2; Length 588;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 QDPQER 9
Db 404 QDPQER 409

RESULT 26
US-09-902-540-16309
; Sequence 16309, Application US/09902540
; Patent No. 6833447
; GENERAL INFORMATION:
; APPLICANT: Goldman, Barry S.
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Wiegand, Roger C.
; TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
; FILE REFERENCE: 38-10(15849)B
; CURRENT APPLICATION NUMBER: US/09/902,540
; CURRENT FILING DATE: 2001-07-10
```

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; PRIOR APPLICATION NUMBER: 60/217,883
; PRIOR FILING DATE: 2000-07-10
; NUMBER OF SEQ ID NOS: 16825
; SEQ ID NO 16309
; LENGTH: 910
; TYPE: PRT
; ORGANISM: Myxococcus xanthus
; US-09-902-540-16309

Query Match
Best Local Similarity 68.8%; Score 33; DB 2; Length 910;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
Db 143 AHFIDPOER 151

RESULT 27
US-09-702-705-1816
; Sequence 1816, Application US/09702705
; Patent No. 6504010
; GENERAL INFORMATION:
; APPLICANT: Wang, Tongtong
; APPLICANT: Bangur, Chaitanya S.
; APPLICANT: Lodes, Michael A.
; APPLICANT: Fanger, Gary
; APPLICANT: Vedvick, Tom
; APPLICANT: Carter, Darlick
; APPLICANT: Retter, Marc
; APPLICANT: Mannion, Jane
; APPLICANT: Fan, Liqun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
; FILE REFERENCE: 210121.478C14
; CURRENT APPLICATION NUMBER: US/09/702,705
; CURRENT FILING DATE: 2000-10-30
; NUMBER OF SEQ ID NOS: 1893
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 1816
; LENGTH: 325
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-09-702-705-1816

Query Match
Best Local Similarity 66.7%; Score 32; DB 2; Length 325;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 157 VFEDPOER 164

RESULT 28
US-09-736-457-1816
; Sequence 1816, Application US/09736457
; Patent No. 6509448
; GENERAL INFORMATION:
; APPLICANT: Wang, Tongtong
; APPLICANT: Bangur, Chaitanya S.
; APPLICANT: Lodes, Michael A.
; APPLICANT: Fanger, Gary
; APPLICANT: Vedvick, Tom
; APPLICANT: Carter, Darlick
; APPLICANT: Retter, Marc
; APPLICANT: Mannion, Jane
; APPLICANT: Fan, Liqun
; APPLICANT: Wang, Aljun
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
; TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
; FILE REFERENCE: 210121.478C15
; CURRENT APPLICATION NUMBER: US/09/736,457
```

/ CURRENT FILING DATE: 2000-12-13
/ NUMBER OF SEQ ID NOS: 1864
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 1816
/ LENGTH: 325
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-09-736-457-1816

Query Match 66.7%; Score 32; DB 2; Length 325;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 157 VFEDPORR 164

RESULT 29
US-09-671-325-1816
/ Sequence 1816, Application US/09671325
/ Patent No. 6667154
/ GENERAL INFORMATION:
/ APPLICANT: Wang, Tongrong
/ APPLICANT: Bangur, Chaitanya S.
/ APPLICANT: Lodes, Michael A.
/ APPLICANT: Fanger, Gary
/ APPLICANT: Vedvick, Tom
/ APPLICANT: Carter, Darrick
/ APPLICANT: Retter, Marc
/ APPLICANT: Mannion, Jane
/ TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
/ TITLE OF INVENTION: DIAGNOSIS OF LUNG CANCER
/ FILE REFERENCE: 210121.478C12
/ CURRENT APPLICATION NUMBER: US/09/671,325
/ CURRENT FILING DATE: 2000-09-26
/ NUMBER OF SEQ ID NOS: 1825
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 1816
/ LENGTH: 325
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-09-671-325-1816

Query Match 66.7%; Score 32; DB 2; Length 325;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 157 VFEDPORR 164

RESULT 30
US-10-017-754-1816
/ Sequence 1816, Application US/10017754
/ Patent No. 6858204
/ GENERAL INFORMATION:
/ APPLICANT: Henderson, Robert A.
/ APPLICANT: Wang, Tongrong
/ APPLICANT: Watanabe, Yoshihiro
/ APPLICANT: Johnson, Jeffrey C.
/ APPLICANT: Retter, Marc W.
/ APPLICANT: Marnerakis, Margarita
/ APPLICANT: Carter, Darrick
/ APPLICANT: Fanger, Gary R.
/ APPLICANT: Vedvick, Thomas S.
/ APPLICANT: Bangur, Chaitanya S.
/ APPLICANT: McNabb, Andria
/ TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
/ TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
/ FILE REFERENCE: 210121.478C18

/ CURRENT APPLICATION NUMBER: US/10/017,754
/ CURRENT FILING DATE: 2001-10-29
/ NUMBER OF SEQ ID NOS: 2004
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 1816
/ LENGTH: 325
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-10-017-754-1816

Query Match 66.7%; Score 32; DB 2; Length 325;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 157 VFEDPORR 164

RESULT 31
US-09-631-863A-2
/ Sequence 2, Application US/09631863A
/ Patent No. 6809179
/ GENERAL INFORMATION:
/ APPLICANT: Boehringer Ingelheim International GmbH
/ TITLE OF INVENTION: Tumor-associated Antigen R11
/ FILE REFERENCE: 12211aa
/ CURRENT APPLICATION NUMBER: US/09/631,863A
/ CURRENT FILING DATE: 2000-08-03
/ NUMBER OF SEQ ID NOS: 102
/ SOFTWARE: PatentIn version 2.1
/ SEQ ID NO 2
/ LENGTH: 401
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-09-631-863A-2

Query Match 66.7%; Score 32; DB 2; Length 401;
Best Local Similarity 62.5%; Pred. No. 2.1e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 233 VFEDPORR 240

RESULT 32
US-09-248-796A-16357
/ Sequence 16357, Application US/09248796A
/ Patent No. 6747137
/ GENERAL INFORMATION:
/ APPLICANT: Keith Weinlock et al
/ TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
/ TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
/ FILE REFERENCE: 107196.132
/ CURRENT APPLICATION NUMBER: US/09/248,796A
/ CURRENT FILING DATE: 1999-02-12
/ PRIOR APPLICATION NUMBER: US 60/074,725
/ PRIOR FILING DATE: 1998-02-13
/ PRIOR APPLICATION NUMBER: US 60/096,409
/ PRIOR FILING DATE: 1998-08-13
/ NUMBER OF SEQ ID NOS: 28208
/ SEQ ID NO 16357
/ LENGTH: 529
/ TYPE: PRT
/ ORGANISM: Candida albicans
US-09-248-796A-16357

Query Match 66.7%; Score 32; DB 2; Length 529;
Best Local Similarity 71.4%; Pred. No. 2.9e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9

DB 240 FQDPPEQ 246

RESULT 33
US-09-489-039A-7923
; Sequence 7923, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:

APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 2709.2004001

CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29

NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 7923

LENGTH: 620
TYPE: PRT
ORGANISM: Klebsiella pneumoniae

US-09-489-039A-7923

Query Match 66.7%; Score 32; DB 2; Length 620;
Best Local Similarity 66.7%; Pred. No. 3.4e+02;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 AMFQDPQER 9
DB 601 AAFNDPQNR 609

RESULT 34
US-09-270-767-41556
; Sequence 41556, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:

APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517

SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 41556
LENGTH: 197

TYPE: PRT
ORGANISM: Drosophila melanogaster
FEATURE:

OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-41556

Query Match 64.6%; Score 31; DB 2; Length 197;
Best Local Similarity 71.4%; Pred. No. 1.6e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPQER 8
DB 41 LMQDPQER 47

RESULT 35
US-07-857-224B-46
; Sequence 46, Application US/07857224B
; Patent No. 5958784
; GENERAL INFORMATION:

APPLICANT: Benner, Steven A.
TITLE OF INVENTION: Predicting Folded Structures of Proteins
NUMBER OF SEQUENCES: 114

CORRESPONDENCE ADDRESS:
ADDRESSEE: Steven A. Benner
STREET: Hadlaubstrasse 151

CITY: Zurich
STATE: none
COUNTRY: Switzerland
ZIP: (note: this is an international post code) CH-8092
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5 inch diskette, 1.4 Mb storage

COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 7.0
SOFTWARE: Microsoft Word

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/857,224B
FILING DATE: 03/25/92
CLASSIFICATION: 436

PRIOR APPLICATION DATA: none
TELECOMMUNICATION INFORMATION:
TELEPHONE: (International) 41 1 632 2830
TELEFAX: (International) 41 1 262 2437
TELEX: none

INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:

LENGTH: 271
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE:

DESCRIPTION: protein
ORIGINAL SOURCE:
ORGANISM: Schizosaccharomyces pombe
FEATURE: Protein kinase; Table 8 Column 52
PUBLICATION INFORMATION:

AUTHORS: Hanke, S. K.
AUTHORS: Quinn, A. M.
AUTHORS: Hunter, T.
TITLE: The protein kinase family
JOURNAL: Science

VOLUME: 241
PAGES: 42-52
DATE: 1988

US-07-857-224B-46

Query Match 64.6%; Score 31; DB 1; Length 271;
Best Local Similarity 55.6%; Pred. No. 2.2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFQDPQER 9
DB 230 AVFEDQGR 238

RESULT 36
US-09-270-767-42416
; Sequence 42416, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:

APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094

CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517

SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 42416
LENGTH: 287

TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-42416

Query Match 64.6%; Score 31; DB 2; Length 287;
Best Local Similarity 55.6%; Pred. No. 2.4e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFQDPQER 9
DB 111 AAFNDPQNR 111

```
Db          70 AVFSDDPK 78

RESULT 37
US-09-252-991A-29703
; Sequence 29703, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 29703
; LENGTH: 324
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-29703

Query Match          64.6%; Score 31; DB 2; Length 324;
Best Local Similarity 75.0%; Pred. No. 2.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY          1 AMFDDPK 8
           |||||
           99 AFDDPK 106

RESULT 38
US-09-252-991A-30165
; Sequence 30165, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 30165
; LENGTH: 338
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-30165

Query Match          64.6%; Score 31; DB 2; Length 338;
Best Local Similarity 55.6%; Pred. No. 2.8e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY          1 AMFDDPK 9
           |||||
           128 AFDDPK 136

RESULT 39
US-09-252-991A-22590
; Sequence 22590, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 22590
; LENGTH: 474
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-22590

Query Match          64.6%; Score 31; DB 2; Length 474;
Best Local Similarity 62.5%; Pred. No. 4e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY          2 MFDDPK 9
           |||||
           183 LFDDPK 190

RESULT 40
US-09-252-991A-19511
; Sequence 19511, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 19511
; LENGTH: 484
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-19511

Query Match          64.6%; Score 31; DB 2; Length 484;
Best Local Similarity 62.5%; Pred. No. 4.1e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY          2 MFDDPK 9
           |||||
           124 LFDDPK 131

RESULT 41
US-09-252-991A-20181
; Sequence 20181, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 20181
; LENGTH: 534
; TYPE: PRT
```

ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-20181

Query Match 64.6%; Score 31; DB 2; Length 534;
Best Local Similarity 66.7%; Pred. No. 4.5e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 306 ADFODPDR 314

RESULT 42
US-10-237-551-121
Sequence 121, Application US/10237551
Patent No. 6821519

GENERAL INFORMATION:
APPLICANT: Day, Craig H.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 121
LENGTH: 545
TYPE: PRT
ORGANISM: HSV-2
US-10-237-551-121

Query Match 64.6%; Score 31; DB 2; Length 545;
Best Local Similarity 55.6%; Pred. No. 4.6e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 481 SLWODPPER 489

RESULT 43
US-10-237-551-157
Sequence 157, Application US/10237551
Patent No. 6821519
GENERAL INFORMATION:
APPLICANT: Day, Craig H.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 157
LENGTH: 545
TYPE: PRT
ORGANISM: HSV2
US-10-237-551-157

Query Match 64.6%; Score 31; DB 2; Length 545;
Best Local Similarity 55.6%; Pred. No. 4.6e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 481 SLWODPPER 489

RESULT 44

US-10-237-551-215
Sequence 215, Application US/10237551
Patent No. 6821519

GENERAL INFORMATION:
APPLICANT: Day, Craig H.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 215
LENGTH: 547
TYPE: PRT
ORGANISM: Herpes Simplex Virus Type 2
US-10-237-551-215

Query Match 64.6%; Score 31; DB 2; Length 547;
Best Local Similarity 55.6%; Pred. No. 4.6e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 483 SLWODPPER 491

RESULT 45
US-10-237-551-216
Sequence 216, Application US/10237551
Patent No. 6821519
GENERAL INFORMATION:
APPLICANT: Day, Craig H.
APPLICANT: Hosken, Nancy A.
APPLICANT: Parsons, Joseph M.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF HERPES SIMPLEX VIRUS INFECTION
FILE REFERENCE: 210121.538C3
CURRENT APPLICATION NUMBER: US/10/237,551
CURRENT FILING DATE: 2002-09-06
NUMBER OF SEQ ID NOS: 254
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 216
LENGTH: 547
TYPE: PRT
ORGANISM: Herpes Simplex Virus Type 2
US-10-237-551-216

Query Match 64.6%; Score 31; DB 2; Length 547;
Best Local Similarity 55.6%; Pred. No. 4.6e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 483 SLWODPPER 491

RESULT 46
US-09-949-016-9511
Sequence 9511, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: C1001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768

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;; PRIOR FILING DATE: 2000-10-03
;; PRIOR APPLICATION NUMBER: 60/231,498
;; PRIOR FILING DATE: 2000-09-08
;; NUMBER OF SEQ ID NOS: 207012
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO: 9511
;; LENGTH: 672
;; TYPE: PRT
;; ORGANISM: Human
US-09-949-016-9511
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Db      221 VFRDPOE 227
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US-09-252-991A-21575
; Sequence 21575, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO: 21575
; LENGTH: 740
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-21575
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Db      228 ILQDPBER 235
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RESULT 48
US-09-328-352-5208
; Sequence 5208, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352
; CURRENT FILING DATE: 1999-06-04
; NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO: 5208
; LENGTH: 818
; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-09-328-352-5208
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Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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Db      152 SIFNDPOE 159
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RESULT 49
US-09-902-540-12961
; Sequence 12961, Application US/09902540
; Patent No. 6833447
; GENERAL INFORMATION:
; APPLICANT: Goldman, Barry S.
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Wiegand, Roger C.
; TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
; FILE REFERENCE: 38-10(15849)B
; CURRENT APPLICATION NUMBER: US/09/902,540
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: 60/217,883
; PRIOR FILING DATE: 2000-07-10
; NUMBER OF SEQ ID NOS: 16825
; SEQ ID NO: 12961
; LENGTH: 909
; TYPE: PRT
; ORGANISM: Myxococcus xanthus
US-09-902-540-12961
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Best Local Similarity 85.7%; Pred. No. 7.9e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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Db      727 FQDPQER 733
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RESULT 50
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; Sequence 16539, Application US/09902540
; Patent No. 6833447
; GENERAL INFORMATION:
; APPLICANT: Goldman, Barry S.
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Wiegand, Roger C.
; TITLE OF INVENTION: Myxococcus xanthus Genome Sequences and Uses Thereof
; FILE REFERENCE: 38-10(15849)B
; CURRENT APPLICATION NUMBER: US/09/902,540
; CURRENT FILING DATE: 2001-07-10
; PRIOR APPLICATION NUMBER: 60/217,883
; PRIOR FILING DATE: 2000-07-10
; NUMBER OF SEQ ID NOS: 16825
; SEQ ID NO: 16539
; LENGTH: 100
; TYPE: PRT
; ORGANISM: Myxococcus xanthus
US-09-902-540-16539
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Best Local Similarity 62.5%; Pred. No. 1.2e+02;
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OM protein - protein search, using SW model

Run on: May 5, 2006, 08:50:57 ; Search time 57 Seconds
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Title: US-08-170-344-1

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Searched: 1867569 seqs, 417829326 residues

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Listing first 1000 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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8	48	100.0	158	6	US-11-021-949-13
9	48	100.0	171	4	US-10-472-724-2
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12	48	100.0	243	6	US-11-072-288-1
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14	48	100.0	266	5	US-09-367-309A-1
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49	34	70.8	456	3	US-09-815-453-8	Sequence 8, App1
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52	34	70.8	792	4	US-10-369-493-4878	Sequence 4878, App1
53	34	70.8	792	4	US-10-369-493-7637	Sequence 7637, App1
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58	33	68.8	257	4	US-10-369-493-7151	Sequence 7151, App1
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65	33	68.8	540	4	US-10-425-114-69537	Sequence 69537, App1
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68	33	68.8	568	5	US-10-746-251-2	Sequence 2, App1
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103	32	66.7	352	4	US-10-424-599-176890	Sequence 176890, A	176	31	64.6	643	4	US-10-282-122A-50408	Sequence 50408, A
104	32	66.7	429	4	US-10-369-493-11702	Sequence 11702, A	177	31	64.6	684	6	US-11-097-143-3426	Sequence 3426, Ap
105	32	66.7	429	4	US-10-369-493-14642	Sequence 14642, A	178	31	64.6	779	4	US-10-296-115-856	Sequence 856, App
106	32	66.7	429	4	US-10-369-493-15104	Sequence 15104, A	179	31	64.6	795	4	US-10-437-963-165267	Sequence 165267, A
107	32	66.7	429	5	US-10-732-923-11520	Sequence 11520, A	180	31	64.6	812	4	US-10-238-075-933	Sequence 933, App
108	32	66.7	438	4	US-10-437-963-102765	Sequence 102765, A	181	31	64.6	882	4	US-10-424-599-218862	Sequence 218862, A
109	32	66.7	438	4	US-10-425-115-273682	Sequence 273682, A	182	31	64.6	957	4	US-10-128-714-3242	Sequence 3242, Ap
110	32	66.7	505	4	US-10-156-761-14310	Sequence 14310, A	183	31	64.6	1048	4	US-10-128-714-8242	Sequence 8242, Ap
111	32	66.7	520	4	US-10-425-115-314145	Sequence 314145, A	184	31	64.6	1099	3	US-09-835-976B-10	Sequence 10, Appl
112	32	66.7	538	4	US-10-425-114-52887	Sequence 52887, A	185	31	64.6	1453	4	US-10-282-122A-45027	Sequence 45027, A
113	32	66.7	708	4	US-10-382-600-29	Sequence 29, Appl	186	31	64.6	1812	3	US-09-775-938A-38	Sequence 38, Appl
114	32	66.7	726	4	US-10-408-765A-2999	Sequence 2999, Ap	187	31	64.6	2103	4	US-10-282-122A-44693	Sequence 44693, A
115	32	66.7	726	5	US-10-723-860-1014	Sequence 1014, Ap	188	31	64.6	4623	6	US-11-097-143-40110	Sequence 40110, A
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117	32	66.7	1413	3	US-09-840-743-8	Sequence 8, Appl.1	190	30	62.5	26	3	US-09-864-761-38435	Sequence 38435, A
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119	32	66.7	1657	5	US-10-760-493-27	Sequence 27, Appl	192	30	62.5	70	4	US-10-425-115-338850	Sequence 328850, A
120	32	66.7	3723	5	US-10-844-715-6	Sequence 6, Appl.1	193	30	62.5	76	5	US-10-450-763-60590	Sequence 60590, A
121	32	66.7	4437	4	US-10-314-637-45	Sequence 45, Appl	194	30	62.5	113	4	US-10-425-115-20100	Sequence 290100, A
122	32	66.7	4437	5	US-10-473-193-45	Sequence 45, Appl	195	30	62.5	114	4	US-10-133-628-20	Sequence 20, Appl
123	32	66.7	4685	4	US-10-156-761-10433	Sequence 10433, A	196	30	62.5	140	4	US-10-133-628-20	Sequence 15, Appl
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125	31	64.6	54	4	US-10-437-963-196343	Sequence 196343, A	198	30	62.5	149	6	US-11-021-949-16	Sequence 428, App
126	31	64.6	86	4	US-10-231-417-520	Sequence 520, App	199	30	62.5	155	4	US-10-074-978A-429	Sequence 428, App
127	31	64.6	104	4	US-10-264-237-2007	Sequence 2007, Ap	200	30	62.5	158	4	US-10-074-978A-428	Sequence 29, Appl
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129	31	64.6	143	4	US-10-767-701-50822	Sequence 50822, A	202	30	62.5	160	6	US-11-021-949-32	Sequence 32, Appl
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135	31	64.6	254	4	US-10-501-282-2094	Sequence 2094, Ap	208	30	62.5	228	4	US-10-437-963-114333	Sequence 114333, A
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137	31	64.6	255	4	US-10-112-944-783	Sequence 783, App	210	30	62.5	233	4	US-10-767-701-39714	Sequence 39714, A
138	31	64.6	259	4	US-10-437-963-125614	Sequence 125614, A	211	30	62.5	252	5	US-10-450-763-30401	Sequence 30401, A
139	31	64.6	259	4	US-10-425-115-287577	Sequence 287577, A	212	30	62.5	252	5	US-10-450-763-30401	Sequence 30401, A
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144	31	64.6	270	4	US-10-425-114-55927	Sequence 55927, A	217	30	62.5	316	5	US-10-501-282-6206	Sequence 6206, Ap
145	31	64.6	270	4	US-10-425-115-246937	Sequence 246937, A	218	30	62.5	325	4	US-10-282-122A-72109	Sequence 72109, A
146	31	64.6	270	4	US-10-425-115-246937	Sequence 246937, A	219	30	62.5	335	4	US-10-724-972A-5091	Sequence 5091, Ap
147	31	64.6	297	4	US-10-425-115-283055	Sequence 283055, A	220	30	62.5	336	4	US-10-282-122A-72545	Sequence 72545, A
148	31	64.6	308	3	US-09-934-455-264	Sequence 264, App	221	30	62.5	343	4	US-10-451-467A-944	Sequence 44, Appl
149	31	64.6	308	4	US-10-180-375-202	Sequence 202, App	222	30	62.5	343	4	US-10-259-194A-244	Sequence 244, App
150	31	64.6	308	4	US-10-225-066A-668	Sequence 668, App	223	30	62.5	346	4	US-10-421-654-56	Sequence 56, Appl
151	31	64.6	308	5	US-10-374-780A-2132	Sequence 2132, Ap	224	30	62.5	346	5	US-10-765-907-56	Sequence 56, Appl
152	31	64.6	308	5	US-10-225-066A-668	Sequence 668, App	225	30	62.5	347	3	US-09-738-626-4237	Sequence 4237, Ap
153	31	64.6	308	5	US-10-225-066A-668	Sequence 668, App	226	30	62.5	347	3	US-10-781-014-20	Sequence 20, Appl
154	31	64.6	373	5	US-10-205-331-22	Sequence 54, Appl	227	30	62.5	351	5	US-10-501-282-6208	Sequence 6208, App
155	31	64.6	405	4	US-10-631-467-1532	Sequence 1532, Ap	228	30	62.5	353	4	US-10-288-252-12	Sequence 12, Appl
156	31	64.6	405	5	US-10-289-148-2	Sequence 2, Appl.1	229	30	62.5	353	4	US-10-398-038-12	Sequence 12, Appl
157	31	64.6	405	5	US-10-618-281-50	Sequence 50, Appl	230	30	62.5	353	5	US-10-975-440-12	Sequence 12, Appl
158	31	64.6	461	4	US-10-437-963-190778	Sequence 190778, A	231	30	62.5	353	5	US-10-501-282-6210	Sequence 6210, Ap
159	31	64.6	470	4	US-10-425-115-243386	Sequence 243386, A	232	30	62.5	384	5	US-10-369-493-11033	Sequence 11033, Ap
160	31	64.6	519	4	US-10-437-963-190810	Sequence 190810, A	233	30	62.5	421	4	US-10-425-115-217184	Sequence 217184, A
161	31	64.6	545	4	US-10-437-963-180705	Sequence 180705, A	234	30	62.5	431	4	US-10-425-115-296546	Sequence 296546, A
162	31	64.6	545	4	US-10-121-988-121	Sequence 121, App	235	30	62.5	434	4	US-10-425-114-68860	Sequence 68860, A
163	31	64.6	545	4	US-10-121-988-121	Sequence 121, App	236	30	62.5	443	5	US-10-450-763-54789	Sequence 54789, A
164	31	64.6	545	4	US-10-200-562-121	Sequence 121, App	237	30	62.5	443	5	US-10-183-687-400	Sequence 400, App
165	31	64.6	545	4	US-10-200-562-121	Sequence 121, App	238	30	62.5	453	4	US-10-282-122A-73183	Sequence 73183, A
166	31	64.6	545	4	US-10-237-551-121	Sequence 121, App	239	30	62.5	466	3	US-09-815-942-14017	Sequence 14017, A
167	31	64.6	545	4	US-10-237-551-157	Sequence 157, App	240	30	62.5	466	4	US-10-282-122A-59513	Sequence 59513, A
168	31	64.6	545	5	US-10-945-050-121	Sequence 12, Appl	241	30	62.5	466	4	US-10-282-122A-76008	Sequence 76008, A
169	31	64.6	545	5	US-10-945-050-121	Sequence 121, App	242	30	62.5	466	4	US-10-425-115-215242	Sequence 215242, A
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171	31	64.6	547	4	US-10-237-551-215	Sequence 215, App	244	30	62.5	491	5	US-10-732-923-1054	Sequence 1054, Ap
172	31	64.6	547	5	US-10-945-050-215	Sequence 215, App	245	30	62.5	491	5	US-10-732-923-1054	Sequence 1054, Ap
173	31	64.6	547	5	US-10-945-050-216	Sequence 216, App	246	30	62.5	492	5	US-10-732-923-1054	Sequence 1054, Ap

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252	30	62.5	513	3	US-10-473-339-4	Sequence 4, App11	325	30	62.5	1246	5	US-10-732-923-17025	Sequence 17025, A
253	30	62.5	516	4	US-10-168-844-8	Sequence 8, App11	326	30	62.5	1246	5	US-10-732-923-17026	Sequence 17026, A
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256	30	62.5	534	5	US-10-450-763-60591	Sequence 60591, A	329	30	62.5	1412	4	US-10-437-963-20158	Sequence 20158,
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258	30	62.5	543	5	US-10-732-923-23525	Sequence 23525, A	331	30	62.5	1479	3	US-10-474-794-305	Sequence 305, App
259	30	62.5	567	5	US-10-450-763-55198	Sequence 55198, A	332	30	62.5	1479	5	US-10-979-159-105	Sequence 305, App
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262	30	62.5	588	5	US-10-732-923-13420	Sequence 13420, A	335	30	62.5	1637	4	US-10-437-963-18037	Sequence 18037, A
263	30	62.5	588	5	US-10-732-923-13422	Sequence 13422, A	336	30	62.5	1661	5	US-10-450-763-55074	Sequence 55074, A
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265	30	62.5	617	4	US-10-437-963-13490	Sequence 143490,	338	30	62.5	2098	4	US-10-132-134-36	Sequence 36, App1
266	30	62.5	659	4	US-10-074-978A-110	Sequence 110, App	339	30	62.5	2109	4	US-10-369-493-6346	Sequence 6346, Ap
267	30	62.5	659	4	US-10-403-161-62	Sequence 62, App1	340	30	62.5	3362	5	US-10-378-083-6	Sequence 8, App11
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270	30	62.5	681	4	US-10-767-701-44840	Sequence 44840, A	343	30	62.5	3798	4	US-10-014-717-6	Sequence 6, App11
271	30	62.5	681	5	US-10-725-329-29	Sequence 29, App1	344	30	62.5	4150	3	US-09-808-880-2	Sequence 23144, A
272	30	62.5	681	5	US-10-737-450-108	Sequence 108, App	345	30	62.5	4273	3	US-10-369-493-23144	Sequence 2, App11
273	30	62.5	681	5	US-10-885-921-10	Sequence 10, App1	346	30	62.5	5215	3	US-09-860-846-2	Sequence 2, App11
274	30	62.5	681	5	US-10-725-121-29	Sequence 29, App1	347	30	62.5	5215	3	US-09-988-344B-2	Sequence 2, App11
275	30	62.5	694	5	US-10-450-763-57994	Sequence 57994, A	348	30	62.5	5215	3	US-09-836-821-2	Sequence 2, App11
276	30	62.5	700	4	US-10-390-585-12	Sequence 12, App1	349	30	62.5	5215	3	US-10-271-889-45	Sequence 45, App1
277	30	62.5	717	4	US-10-054-044A-2	Sequence 2, App11	350	30	62.5	6145	4	US-10-156-761-7962	Sequence 7962, Ap
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280	30	62.5	792	4	US-10-087-192-1977	Sequence 1977, Ap	353	30	62.5	7257	5	US-10-732-923-20620	Sequence 20620, A
281	30	62.5	796	4	US-10-369-493-703	Sequence 703, App	354	30	62.5	7257	5	US-10-314-657-20621	Sequence 20621, A
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283	30	62.5	913	5	US-10-450-763-38474	Sequence 38474, A	356	30	62.5	7349	5	US-10-424-599-22666	Sequence 22666, A
284	30	62.5	913	5	US-10-295-027-490	Sequence 490, App	357	30	62.5	8026	4	US-10-424-599-22666	Sequence 22666, A
285	30	62.5	936	4	US-10-173-999-18	Sequence 18, App1	358	30	62.5	8026	4	US-10-424-599-22666	Sequence 22666, A
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287	30	62.5	959	5	US-10-840-512-222	Sequence 222, App	360	30	62.5	9477	9	US-10-203-295-37	Sequence 37, App1
288	30	62.5	962	4	US-10-437-963-145906	Sequence 145906,	361	29	60.4	9	4	US-10-033-662-8	Sequence 8, App1
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290	30	62.5	967	4	US-10-173-999-16	Sequence 16, App1	363	29	60.4	50	5	US-10-776-013-271	Sequence 271, App
291	30	62.5	993	4	US-10-173-999-14	Sequence 14, App1	364	29	60.4	54	5	US-10-450-763-56421	Sequence 56421, A
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293	30	62.5	1000	4	US-10-173-999-22	Sequence 22, App1	366	29	60.4	66	4	US-10-424-599-217760	Sequence 217760,
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298	30	62.5	1014	4	US-10-225-567A-428	Sequence 428, App	371	29	60.4	84	4	US-10-424-599-223283	Sequence 223283,
299	30	62.5	1014	4	US-10-295-027-492	Sequence 492, App	372	29	60.4	89	4	US-10-424-599-212193	Sequence 212193,
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307	30	62.5	1014	6	US-11-070-456-139	Sequence 139, App	380	29	60.4	101	5	US-10-450-763-55120	Sequence 55120, A
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311	30	62.5	1073	6	US-11-093-888-14	Sequence 14, App1	384	29	60.4	110	4	US-10-424-599-229328	Sequence 229328, A
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313	30	62.5	1085	4	US-10-174-363-12	Sequence 12, App1	386	29	60.4	112	4	US-10-205-882-154	Sequence 154, App
314	30	62.5	1085	6	US-11-093-888-12	Sequence 12, App1	387	29	60.4	112	4	US-10-425-115-216644	Sequence 216644, A
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316	30	62.5	1140	4	US-10-468-406-8	Sequence 8, App11	389	29	60.4	114	4	US-10-767-701-42264	Sequence 42264, A
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319	30	62.5	1245	5	US-10-732-923-17019	Sequence 17019, A	392	29	60.4	125	4	US-10-424-599-212194	Sequence 212194, A

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394	29	60.4	126	4	US-10-425-115-221355	Sequence 221355,	467	29	60.4	267	4	US-10-424-599-69756	Sequence 269756,
395	29	60.4	130	4	US-10-424-599-146022	Sequence 146022,	468	29	60.4	269	4	US-10-369-493-8813	Sequence 2813, Ap
396	29	60.4	130	4	US-10-424-599-146379	Sequence 146379,	469	29	60.4	274	4	US-10-218-654-53	Sequence 53, Appl
397	29	60.4	135	4	US-10-220-120-300	Sequence 300, App	470	29	60.4	274	4	US-10-262-439-53	Sequence 53, Appl
398	29	60.4	145	4	US-10-425-115-284765	Sequence 284765,	471	29	60.4	283	4	US-10-374-780A-2012	Sequence 2012, Ap
399	29	60.4	148	6	US-11-021-949-17	Sequence 17, Appl	472	29	60.4	283	4	US-10-412-659B-312	Sequence 312, App
400	29	60.4	149	6	US-11-021-949-18	Sequence 18, Appl	473	29	60.4	295	4	US-10-425-114-53449	Sequence 53449, A
401	29	60.4	151	6	US-11-021-949-24	Sequence 24, Appl	474	29	60.4	296	4	US-10-425-114-64864	Sequence 64864, A
402	29	60.4	151	6	US-11-021-949-24	Sequence 24, Appl	475	29	60.4	297	6	US-11-097-143-11809	Sequence 31809, A
403	29	60.4	153	6	US-10-424-599-217664	Sequence 217664,	476	29	60.4	298	4	US-10-389-566-565	Sequence 566, App
404	29	60.4	155	6	US-11-021-949-22	Sequence 22, Appl	477	29	60.4	302	4	US-10-425-115-289788	Sequence 289788,
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409	29	60.4	161	4	US-10-767-701-41214	Sequence 41214, A	482	29	60.4	320	4	US-10-437-963-123469	Sequence 123469,
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411	29	60.4	165	4	US-10-363-829-269	Sequence 269, App	484	29	60.4	321	4	US-10-425-114-58590	Sequence 58590, A
412	29	60.4	168	4	US-10-767-701-56173	Sequence 56173, A	485	29	60.4	329	5	US-10-474-792-556	Sequence 556, App
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419	29	60.4	189	4	US-10-424-599-193365	Sequence 193365,	492	29	60.4	353	5	US-10-732-923-4430	Sequence 4430, Ap
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422	29	60.4	192	4	US-10-425-115-236290	Sequence 236290,	495	29	60.4	356	4	US-10-282-122A-75372	Sequence 140, App
423	29	60.4	192	4	US-10-425-115-236292	Sequence 236292,	496	29	60.4	356	4	US-10-210-112-148	Sequence 74, Appl
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425	29	60.4	192	4	US-10-425-115-236301	Sequence 236301,	498	29	60.4	356	5	US-10-471-115-9	Sequence 74, Appl
426	29	60.4	192	4	US-10-425-115-236303	Sequence 236303,	499	29	60.4	356	5	US-10-980-387-74	Sequence 471, App
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428	29	60.4	193	4	US-10-437-963-130350	Sequence 130350,	501	29	60.4	360	5	US-09-925-763-1068	Sequence 1068, Ap
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432	29	60.4	195	4	US-10-156-761-13170	Sequence 13170, A	505	29	60.4	370	4	US-10-451-168-73	Sequence 73, Appl
433	29	60.4	195	4	US-10-425-114-51080	Sequence 51080, A	506	29	60.4	370	5	US-10-980-387-73	Sequence 152, App
434	29	60.4	196	5	US-10-450-763-58941	Sequence 58941, A	507	29	60.4	373	4	US-10-205-823-152	Sequence 72, Appl
435	29	60.4	198	4	US-10-425-114-47381	Sequence 47381, A	508	29	60.4	373	4	US-10-287-226-72	Sequence 72, Appl
436	29	60.4	200	4	US-10-437-963-141520	Sequence 141520,	509	29	60.4	373	4	US-10-287-226-76	Sequence 76, Appl
437	29	60.4	200	4	US-10-424-599-186185	Sequence 186185,	510	29	60.4	373	5	US-10-792-571-18	Sequence 797, App
438	29	60.4	209	4	US-10-425-114-59374	Sequence 59374, A	511	29	60.4	373	5	US-10-631-461-797	Sequence 152, App
439	29	60.4	212	4	US-10-425-114-63052	Sequence 63052, A	512	29	60.4	373	6	US-11-051-454-152	Sequence 1658, Ap
440	29	60.4	213	4	US-10-032-585-7714	Sequence 7714, Ap	513	29	60.4	379	4	US-10-369-493-6658	Sequence 6658, Ap
441	29	60.4	213	4	US-10-425-114-47712	Sequence 47712, A	514	29	60.4	383	4	US-10-437-963-162754	Sequence 162754,
442	29	60.4	213	4	US-10-425-114-60506	Sequence 60506, A	515	29	60.4	387	4	US-10-287-226-70	Sequence 70, Appl
443	29	60.4	215	4	US-10-425-114-59760	Sequence 59760, A	516	29	60.4	389	4	US-10-156-761-11356	Sequence 11356, A
444	29	60.4	216	4	US-10-425-114-59455	Sequence 59455, A	517	29	60.4	391	4	US-10-424-599-13328	Sequence 564, App
445	29	60.4	216	4	US-10-425-114-48281	Sequence 48281, A	518	29	60.4	392	4	US-10-389-566-564	Sequence 564, App
446	29	60.4	217	4	US-10-425-114-59358	Sequence 59358, A	519	29	60.4	396	4	US-10-104-047-2919	Sequence 2919, App
447	29	60.4	217	4	US-10-425-114-59618	Sequence 59618, A	520	29	60.4	397	4	US-10-732-923-11576	Sequence 24144, A
448	29	60.4	217	4	US-10-425-114-59618	Sequence 59618, A	521	29	60.4	407	5	US-10-425-115-354223	Sequence 354223,
449	29	60.4	217	4	US-10-425-114-59642	Sequence 59642, A	522	29	60.4	415	4	US-10-425-115-354223	Sequence 59107, A
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452	29	60.4	229	4	US-10-104-047-3781	Sequence 3781, Ap	525	29	60.4	429	5	US-10-732-923-11685	Sequence 11685, A
453	29	60.4	231	4	US-10-425-115-343579	Sequence 343579,	526	29	60.4	429	5	US-10-732-923-11685	Sequence 11685, A
454	29	60.4	233	4	US-10-425-114-41550	Sequence 41550, A	527	29	60.4	430	5	US-10-732-923-11576	Sequence 11576, A
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457	29	60.4	247	5	US-10-882-122A-69667	Sequence 69667, A	530	29	60.4	433	4	US-10-425-114-72358	Sequence 20657, A
458	29	60.4	251	5	US-10-732-923-1573	Sequence 1573, Ap	531	29	60.4	434	4	US-10-369-493-20657	Sequence 11421, A
459	29	60.4	254	4	US-10-032-585-7420	Sequence 7420, Ap	532	29	60.4	434	5	US-10-732-923-11401	Sequence 11606, A
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461	29	60.4	252	5	US-10-472-928-2832	Sequence 2832, Ap	534	29	60.4	436	4	US-10-425-115-21387	Sequence 11609, A
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463	29	60.4	255	4	US-10-562-439-58	Sequence 58, Appl	536	29	60.4	441	4	US-10-347-470A-6	Sequence 17646, A
464	29	60.4	256	5	US-10-424-599-252228	Sequence 252228,	537	29	60.4	441	4	US-10-347-470A-6	Sequence 17646, A
465	29	60.4	267	4	US-10-369-493-2346	Sequence 2346, Ap	538	29	60.4	441	6	US-11-097-143-17646	Sequence 17646, A

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510	29	60.4	454	3	US-09-739-254-165	Sequence 165, App	613	29	60.4	547	4	US-10-243-425-40	Sequence 40, Appl
541	29	60.4	454	3	US-09-904-615-165	Sequence 165, App	614	29	60.4	547	4	US-10-243-446-60	Sequence 40, Appl
542	29	60.4	454	3	US-10-054-988-165	Sequence 165, App	615	29	60.4	547	4	US-10-245-874-40	Sequence 40, Appl
543	29	60.4	457	4	US-10-195-518-6	Sequence 6, Appl	616	29	60.4	547	4	US-10-242-653-40	Sequence 40, Appl
544	29	60.4	465	4	US-10-282-122A-55825	Sequence 55825, A	617	29	60.4	547	4	US-10-243-167-40	Sequence 40, Appl
545	29	60.4	466	3	US-09-813-242-10021	Sequence 10021, A	618	29	60.4	547	4	US-10-243-388-40	Sequence 40, Appl
546	29	60.4	466	3	US-10-282-122A-56405	Sequence 56405, A	619	29	60.4	547	4	US-10-244-947-40	Sequence 40, Appl
547	29	60.4	466	4	US-10-425-114-46206	Sequence 46206, A	620	29	60.4	547	4	US-10-244-968-40	Sequence 40, Appl
548	29	60.4	469	4	US-10-425-115-321155	Sequence 321155,	621	29	60.4	547	4	US-10-244-990-40	Sequence 40, Appl
549	29	60.4	478	4	US-10-424-599-272789	Sequence 272789,	622	29	60.4	547	4	US-10-245-079-40	Sequence 40, Appl
550	29	60.4	478	5	US-10-732-923-24143	Sequence 24143, A	623	29	60.4	547	4	US-10-245-127-40	Sequence 40, Appl
551	29	60.4	480	4	US-10-009-823A-11	Sequence 11, Appl	624	29	60.4	547	4	US-10-245-207-40	Sequence 40, Appl
552	29	60.4	484	4	US-10-425-114-40095	Sequence 40095, A	625	29	60.4	547	4	US-10-245-646-40	Sequence 40, Appl
553	29	60.4	499	4	US-10-156-761-11491	Sequence 11491, A	626	29	60.4	547	4	US-10-245-665-40	Sequence 40, Appl
554	29	60.4	500	4	US-10-369-493-22495	Sequence 22495, A	627	29	60.4	547	4	US-10-245-699-40	Sequence 40, Appl
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556	29	60.4	506	4	US-10-094-749-2229	Sequence 2229, Ap	629	29	60.4	547	4	US-10-245-890-40	Sequence 40, Appl
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558	29	60.4	517	6	US-11-097-143-1128	Sequence 1128, Ap	631	29	60.4	547	4	US-10-245-900-40	Sequence 40, Appl
559	29	60.4	518	4	US-10-437-963-195842	Sequence 195842,	632	29	60.4	547	4	US-10-247-058-40	Sequence 40, Appl
560	29	60.4	520	5	US-10-732-923-10663	Sequence 10663, A	633	29	60.4	547	4	US-10-247-454-40	Sequence 40, Appl
561	29	60.4	530	4	US-10-425-115-289786	Sequence 289786,	634	29	60.4	547	4	US-10-247-491-40	Sequence 40, Appl
562	29	60.4	531	4	US-10-425-114-70256	Sequence 70256, A	635	29	60.4	547	4	US-10-238-261-40	Sequence 40, Appl
563	29	60.4	532	5	US-10-732-923-13633	Sequence 13633, A	636	29	60.4	547	4	US-10-238-324-40	Sequence 40, Appl
564	29	60.4	532	5	US-10-732-923-13635	Sequence 13635, A	637	29	60.4	547	4	US-10-238-324-40	Sequence 40, Appl
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567	29	60.4	536	3	US-09-815-242-11857	Sequence 11857, A	640	29	60.4	547	4	US-10-242-652-40	Sequence 40, Appl
568	29	60.4	537	4	US-10-425-114-70196	Sequence 70196, A	641	29	60.4	547	4	US-10-242-990-40	Sequence 40, Appl
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573	29	60.4	542	4	US-10-437-963-148123	Sequence 148123,	646	29	60.4	547	4	US-10-243-499-40	Sequence 40, Appl
574	29	60.4	546	4	US-10-425-115-245497	Sequence 245497,	647	29	60.4	547	4	US-10-244-995-40	Sequence 40, Appl
575	29	60.4	547	4	US-10-445-752-40	Sequence 40, Appl	648	29	60.4	547	4	US-10-244-995-40	Sequence 40, Appl
576	29	60.4	547	4	US-10-445-859-40	Sequence 40, Appl	649	29	60.4	547	4	US-10-245-230-40	Sequence 40, Appl
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578	29	60.4	547	4	US-10-445-107-40	Sequence 40, Appl	651	29	60.4	547	4	US-10-245-479-40	Sequence 40, Appl
579	29	60.4	547	4	US-10-445-143-40	Sequence 40, Appl	652	29	60.4	547	4	US-10-245-499-40	Sequence 40, Appl
580	29	60.4	547	4	US-10-245-771-40	Sequence 40, Appl	653	29	60.4	547	4	US-10-245-812-40	Sequence 40, Appl
581	29	60.4	547	4	US-10-245-851-40	Sequence 40, Appl	654	29	60.4	547	4	US-10-245-812-40	Sequence 40, Appl
582	29	60.4	547	4	US-10-245-883-40	Sequence 40, Appl	655	29	60.4	547	4	US-10-245-852-40	Sequence 40, Appl
583	29	60.4	547	4	US-10-237-535-40	Sequence 40, Appl	656	29	60.4	547	4	US-10-245-881-40	Sequence 40, Appl
584	29	60.4	547	4	US-10-338-183-40	Sequence 40, Appl	657	29	60.4	547	4	US-10-245-875-40	Sequence 40, Appl
585	29	60.4	547	4	US-10-338-283-40	Sequence 40, Appl	658	29	60.4	547	4	US-10-245-913-40	Sequence 40, Appl
586	29	60.4	547	4	US-10-338-370-40	Sequence 40, Appl	659	29	60.4	547	4	US-10-245-913-40	Sequence 40, Appl
587	29	60.4	547	4	US-10-345-055-40	Sequence 40, Appl	660	29	60.4	547	4	US-10-245-910-40	Sequence 40, Appl
588	29	60.4	547	4	US-10-345-147-40	Sequence 40, Appl	661	29	60.4	547	4	US-10-246-080-40	Sequence 40, Appl
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594	29	60.4	547	4	US-10-243-409-40	Sequence 40, Appl	667	29	60.4	547	4	US-10-245-810-40	Sequence 40, Appl
595	29	60.4	547	4	US-10-245-621-40	Sequence 40, Appl	668	29	60.4	547	4	US-10-245-910-40	Sequence 40, Appl
596	29	60.4	547	4	US-10-245-880-40	Sequence 40, Appl	669	29	60.4	547	4	US-10-246-098-40	Sequence 40, Appl
597	29	60.4	547	4	US-10-245-033-40	Sequence 40, Appl	670	29	60.4	547	4	US-10-237-496-40	Sequence 40, Appl
598	29	60.4	547	4	US-10-243-093-40	Sequence 40, Appl	671	29	60.4	547	4	US-10-242-074-40	Sequence 40, Appl
599	29	60.4	547	4	US-10-245-185-40	Sequence 40, Appl	672	29	60.4	547	4	US-10-242-505-40	Sequence 40, Appl
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601	29	60.4	547	4	US-10-245-473-40	Sequence 40, Appl	674	29	60.4	547	4	US-10-243-282-40	Sequence 40, Appl
602	29	60.4	547	4	US-10-245-770-40	Sequence 40, Appl	675	29	60.4	547	4	US-10-243-402-40	Sequence 40, Appl
603	29	60.4	547	4	US-10-245-877-40	Sequence 40, Appl	676	29	60.4	547	4	US-10-243-431-40	Sequence 40, Appl
604	29	60.4	547	4	US-10-246-976-40	Sequence 40, Appl	677	29	60.4	547	4	US-10-245-164-40	Sequence 40, Appl
605	29	60.4	547	4	US-10-243-320-40	Sequence 40, Appl	678	29	60.4	547	4	US-10-210-951-68	Sequence 68, Appl
606	29	60.4	547	4	US-10-242-743-40	Sequence 40, Appl	679	29	60.4	547	4	US-10-244-972-40	Sequence 40, Appl
607	29	60.4	547	4	US-10-242-845-40	Sequence 40, Appl	680	29	60.4	547	4	US-10-244-972-40	Sequence 40, Appl
608	29	60.4	547	4	US-10-237-636-40	Sequence 40, Appl	681	29	60.4	547	4	US-10-197-942-40	Sequence 40, Appl
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610	29	60.4	547	4	US-10-238-346-40	Sequence 40, Appl	683	29	60.4	547	4	US-10-238-196-40	Sequence 40, Appl
611	29	60.4	547	4	US-10-238-411-40	Sequence 40, Appl	684	29	60.4	547	4	US-10-245-013-40	Sequence 40, Appl

685	29	60.4	547	4	US-10-211-858-68	Sequence 68, Appl	758	29	60.4	1346	4	US-10-271-889-37	Sequence 37, Appl
686	29	60.4	549	4	US-10-259-199A-258	Sequence 258, App	759	29	60.4	1346	5	US-10-468-828-8	Sequence 4, Appl1
687	29	60.4	552	4	US-10-369-493-20174	Sequence 20174, A	760	29	60.4	1346	5	US-10-846-335-4	Sequence 4, Appl1
688	29	60.4	555	3	US-09-765-205-16	Sequence 16, Appl	761	29	60.4	1353	5	US-10-732-923-12887	Sequence 12887, A
689	29	60.4	555	5	US-10-347-669-16	Sequence 16, Appl	762	29	60.4	1388	5	US-10-733-923-18237	Sequence 18237, A
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691	29	60.4	557	4	US-10-128-714-3133	Sequence 3133, Ap	764	29	60.4	1478	4	US-10-369-493-17588	Sequence 17588, A
692	29	60.4	557	4	US-10-369-493-19143	Sequence 19143, A	765	29	60.4	1482	3	US-09-940-316B-21	Sequence 21, Appl
693	29	60.4	558	5	US-10-450-763-58946	Sequence 58946, A	766	29	60.4	1488	3	US-09-940-316B-17	Sequence 17, Appl
694	29	60.4	559	4	US-10-156-761-10881	Sequence 10881, A	767	29	60.4	1509	3	US-09-940-316B-23	Sequence 23, Appl
695	29	60.4	560	4	US-10-389-647-654	Sequence 654, App	768	29	60.4	1517	3	US-09-940-316B-19	Sequence 19, Appl
696	29	60.4	569	6	US-11-097-143-20880	Sequence 20880, A	769	29	60.4	1555	6	US-11-097-143-3333	Sequence 3333, Ap
697	29	60.4	569	6	US-11-097-143-28650	Sequence 28650, A	770	29	60.4	1557	3	US-09-940-316B-27	Sequence 27, Appl
698	29	60.4	577	5	US-10-732-923-10712	Sequence 10712, A	771	29	60.4	1562	3	US-09-861-289-35	Sequence 35, Appl
699	29	60.4	578	4	US-10-425-114-55229	Sequence 55229, A	772	29	60.4	1562	3	US-09-861-289-35	Sequence 35, Appl
700	29	60.4	582	4	US-10-424-599-268404	Sequence 268404, A	773	29	60.4	1562	3	US-09-988-384B-35	Sequence 35, Appl
701	29	60.4	594	4	US-10-425-115-187150	Sequence 187150, A	774	29	60.4	1562	3	US-09-836-821-35	Sequence 35, Appl
702	29	60.4	595	4	US-10-425-115-289787	Sequence 289787, A	775	29	60.4	1562	3	US-09-793-708-3	Sequence 3, Appl1
703	29	60.4	599	4	US-10-282-123A-63567	Sequence 63567, A	776	29	60.4	1562	4	US-10-201-365-4	Sequence 4, Appl1
704	29	60.4	609	5	US-10-213-974-25	Sequence 25, Appl	777	29	60.4	1562	4	US-10-160-539-3	Sequence 3, Appl1
705	29	60.4	612	4	US-10-437-963-185375	Sequence 185375, A	778	29	60.4	1562	4	US-10-271-889-35	Sequence 35, Appl
706	29	60.4	640	4	US-10-425-114-37496	Sequence 37496, A	779	29	60.4	1562	5	US-10-468-828-3	Sequence 3, Appl1
707	29	60.4	641	4	US-10-437-963-188700	Sequence 188700, A	780	29	60.4	1562	5	US-10-846-335-3	Sequence 3, Appl1
708	29	60.4	643	4	US-10-424-599-159882	Sequence 159882, A	781	29	60.4	1568	4	US-10-228-148B-5	Sequence 5, Appl1
709	29	60.4	666	4	US-10-437-963-197206	Sequence 197206, A	782	29	60.4	1574	3	US-09-940-316B-25	Sequence 25, Appl
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711	29	60.4	686	4	US-10-425-115-213836	Sequence 213836, A	784	29	60.4	1578	3	US-09-940-316B-31	Sequence 31, Appl
712	29	60.4	697	5	US-10-450-763-53766	Sequence 53766, A	785	29	60.4	1588	3	US-09-940-316B-29	Sequence 29, Appl
713	29	60.4	708	5	US-10-450-763-50466	Sequence 50466, A	786	29	60.4	1589	5	US-10-450-763-45556	Sequence 34556, A
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715	29	60.4	735	4	US-10-408-765A-2988	Sequence 2988, Ap	788	29	60.4	1589	5	US-10-450-763-52962	Sequence 52962, A
716	29	60.4	750	4	US-10-437-963-148541	Sequence 148541, A	789	29	60.4	1589	5	US-10-450-763-88863	Sequence 88863, A
717	29	60.4	757	5	US-10-450-763-31940	Sequence 31940, A	790	29	60.4	1589	5	US-10-450-763-60458	Sequence 60458, A
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ALIGNMENTS

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; Publication No. US20030103964A1
; GENERAL INFORMATION:
; APPLICANT: Delibai, Charles
; APPLICANT: Berzofsky, Jay
; APPLICANT: Gulukota, Kamalakara
; APPLICANT: Vaccaro, Dennis
; APPLICANT: Weng, Zhiping
; APPLICANT: Zhang, Chao
; TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
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; CURRENT FILING DATE: 2002-04-26
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US-10-133-210-277
Query Match 100.0%; Score 48; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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DB 1 AMFODPOER 9
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RESULT 2
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; Sequence 81, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
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; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
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US-10-751-845-81
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```

```
QY 1 AMFODPOER 9
DB 1 AMFODPOER 9
```

```
RESULT 3
US-10-751-845-64
; Sequence 64, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 64
; LENGTH: 20
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-64
Query Match 100.0%; Score 48; DB 5; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.054;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
DB 1 AMFODPOER 9
```

```
RESULT 4
US-10-476-570-59
; Sequence 59, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUELLE-MORAVILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
```

```

; TITLE OF INVENTION: Papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 59
; LENGTH: 22
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 1-22
US-10-476-570-59
```

```
Query Match          100.0%; Score 48; DB 4; Length 22;
Best Local Similarity 100.0%; Pred. No. 0.06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
    |||||
Db 7 AMFODPOER 15
```

```

RESULT 5
US-10-751-845-126
; Sequence 126, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 126
; LENGTH: 117
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-126
```

```
Query Match          100.0%; Score 48; DB 5; Length 117;
Best Local Similarity 100.0%; Pred. No. 0.34;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
    |||||
Db 1 AMFODPOER 9
```

```

RESULT 6
US-10-858-384-2
; Sequence 2, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
```

```

; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-10-858-384-2
```

```
Query Match          100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.46;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
    |||||
Db 7 AMFODPOER 15
```

```

RESULT 7
US-10-367-057-16
; Sequence 16, Application US/10367057
; Publication No. US2005010054A1
; GENERAL INFORMATION:
; APPLICANT: Cutbill, Scott;
; APPLICANT: Jackson, Amanda;
; APPLICANT: Lewin, David A.;
; APPLICANT: Ooi, Chean Eng
; TITLE OF INVENTION: Complexes and Methods of Using Same
; FILE REFERENCE: 21402-559
; CURRENT APPLICATION NUMBER: US/10/367,057
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: 60/256,911
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 198
; SOFTWARE: CutsSeq version 0.1
; SEQ ID NO 16
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-367-057-16
```

```
Query Match          100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.46;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
    |||||
Db 7 AMFODPOER 15
```

```

RESULT 8
US-11-021-949-13
; Sequence 13, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SAMIENTO, CHAMORRO SONOZA
; APPLICANT: SCHWEITZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
```


;; PRIOR APPLICATION NUMBER: 60/532,373
;; PRIOR FILING DATE: 2003-12-23
;; NUMBER OF SEQ ID NOS: 361
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 13
;; LENGTH: 158
;; TYPE: PRT
;; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match 100.0%; Score 48; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.46;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||||
DB 7 AMFODPOER 15

RESULT 9
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; PRIOR FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03271
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match 100.0%; Score 48; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 0.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||||
DB 12 AMFODPOER 20

RESULT 10
US-10-751-845-157
; Sequence 157, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; PRIOR FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0

;; SEQ ID NO 157
;; LENGTH: 236
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-157

Query Match 100.0%; Score 48; DB 5; Length 236;
Best Local Similarity 100.0%; Pred. No. 0.63;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||||
DB 1 AMFODPOER 9

RESULT 11
US-10-751-845-158
; Sequence 158, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; PRIOR FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 158
; LENGTH: 237
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-158

Query Match 100.0%; Score 48; DB 5; Length 237;
Best Local Similarity 100.0%; Pred. No. 0.7;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
|||||||
DB 2 AMFODPOER 10

RESULT 12
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOU, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; PRIOR FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18

NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-11-072-288-1

Query Match 100.0%; Score 48; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.71;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 35 AMFODPOER 43

RESULT 13
US-10-751-845-160
Sequence 160, Application US/10751845
Publication No. US20050100928A1
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Urban, Robert G.
APPLICANT: Chic, Roman M.
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
FILE REFERENCE: 08191-013001
CURRENT APPLICATION NUMBER: US/10/751,845
PRIOR FILING DATE: 2004-01-05
PRIOR APPLICATION NUMBER: US/09/664,225
PRIOR FILING DATE: 2000-08-18
PRIOR APPLICATION NUMBER: US 60/169,846
PRIOR FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: US 60/154,665
PRIOR FILING DATE: 1999-09-16
NUMBER OF SEQ ID NOS: 163
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 160
LENGTH: 261
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-160

Query Match 100.0%; Score 48; DB 5; Length 261;
Best Local Similarity 100.0%; Pred. No. 0.77;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 26 AMFODPOER 34

RESULT 14
US-09-367-309A-1
Sequence 1, Application US/09367309A
Publication No. US20020081329A1
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU96/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19

NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 48; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.78;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 7 AMFODPOER 15

RESULT 15
US-10-000-903-4
Sequence 4, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Bernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencich, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
PRIOR FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-4

Query Match 100.0%; Score 48; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.81;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 113 AMFODPOER 121

RESULT 16
US-10-899-771-4
Sequence 4, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
PRIOR FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4

```

; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4
```

```
Query Match          100.0%; Score 48; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.81;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
DB 113 AMFODPOER 121
```

```
RESULT 17
US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-10
```

```
Query Match          100.0%; Score 48; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.86;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
DB 132 AMFODPOER 140
```

```
RESULT 18
US-10-899-771-10
; Sequence 10, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
```

```

; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-10
```

```
Query Match          100.0%; Score 48; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.86;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
DB 132 AMFODPOER 140
```

```
RESULT 19
US-10-000-903-6
; Sequence 6, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-6
```

```
Query Match          100.0%; Score 48; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 AMFODPOER 9
DB 113 AMFODPOER 121
```

```
RESULT 20
US-10-899-771-6
; Sequence 6, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
```

SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimeraic protein (protein D from Haemophilus
OTHER INFORMATION: Influenzae B and E6B7 fusion from Human papilloma
OTHER INFORMATION: Virus type 16)
US-10-899-771-6

Query Match 100.0%; Score 48; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 113 AMFODPOER 121

RESULT 21
US-10-000-903-14
Sequence 14, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Bernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Benchelkn, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-14

Query Match 100.0%; Score 48; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 1.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 132 AMFODPOER 140

RESULT 22
US-10-899-771-14
Sequence 14, Application US/10899771
Publication No. US2005001638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24

NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimeraic protein (Clyta from Streptococcus
OTHER INFORMATION: pneumoniae and E6B7 fusion from Human papilloma
OTHER INFORMATION: Virus type 16)
US-10-899-771-14

Query Match 100.0%; Score 48; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 1.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 132 AMFODPOER 140

RESULT 23
US-10-367-095-10
Sequence 10, Application US/10367095
Publication No. US20030228696A1
GENERAL INFORMATION:
APPLICANT: Robin A. Robinson
TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
FILE REFERENCE: 44149-1US1
CURRENT APPLICATION NUMBER: US/10/367,095
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,161
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,118
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,133
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,157
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,156
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,123
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,113
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,154
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,135
PRIOR FILING DATE: 2002-02-14
Remaining Prior Application data removed - See file Wrapper or PALM.
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 10
LENGTH: 536
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-367-095-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 AMFODPOER 9
Db 477 AMFODPOER 485

RESULT 24
US-10-368-046-10

```
; Sequence 10, Application US/10368046
; Publication No. US20040063188A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; APPLICANT: Victoria Cloce
; TITLE OF INVENTION: Method for Isolation and Purification of
; TITLE OF INVENTION: Expressed Gene Products in Vitro
; FILE REFERENCE: 44149-30S1
; CURRENT APPLICATION NUMBER: US/10/368,046
; PRIOR FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-368-046-10

Query Match          100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 AMFODPOER 9
        |||||
Db      477 AMFODPOER 485

RESULT 25
US-10-367-367-10
; Sequence 10, Application US/10367367
; Publication No. US20040121465A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 44149-2US1
; CURRENT APPLICATION NUMBER: US/10/367,367
; PRIOR FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-367-367-10

Query Match          100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 AMFODPOER 9
        |||||
Db      477 AMFODPOER 485

RESULT 27
US-10-177-390-6
; Sequence 6, Application US/10177390
```

```
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-367-367-10

Query Match          100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 AMFODPOER 9
        |||||
Db      477 AMFODPOER 485

RESULT 26
US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:
; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 19065/2132
; CURRENT APPLICATION NUMBER: US/10/918,337
; PRIOR FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-918-337-10

Query Match          100.0%; Score 48; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 AMFODPOER 9
        |||||
Db      477 AMFODPOER 485

RESULT 27
US-10-177-390-6
; Sequence 6, Application US/10177390
```

```
Publication No. US20030143743A1
GENERAL INFORMATION:
APPLICANT: Schuler, Gerold
APPLICANT: N.V. Antwerp Innovatiecentrum
TITLE OF INVENTION: Improved transfection of Eucaryotic Cells with Linear
FILE REFERENCE: 021505wo/JH/ml
CURRENT APPLICATION NUMBER: US/10/177,390
CURRENT FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match          91.7%; Score 44; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 1 MFQDPOER 8

RESULT 28
US-10-484-063-20
Sequence 20, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILTERMO
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 20
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-484-063-20

Query Match          91.7%; Score 44; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 1 MFQDPOER 8

RESULT 29
US-10-484-063-27
Sequence 27, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILTERMO
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
CURRENT FILING DATE: 2004-01-16
```

```
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match          91.7%; Score 44; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 1 MFQDPOER 8

RESULT 30
US-10-118-047-2
Sequence 2, Application US/10118047
Publication No. US20030170835A1
GENERAL INFORMATION:
APPLICANT: CHUNG, BONG HYUN
APPLICANT: LEE, EUN GYO
APPLICANT: HAHM, MOON SUN
APPLICANT: RYU, YEON WOO
APPLICANT: LEE, HAN SEUNG
TITLE OF INVENTION: OPTICALLY ACTIVE ARYL PROPIONIC ACIDS AND USING THE
FILE REFERENCE: 05823.0218-00000
CURRENT APPLICATION NUMBER: US/10/118,047
CURRENT FILING DATE: 2002-07-03
PRIOR APPLICATION NUMBER: KR 2002-2809
PRIOR FILING DATE: 2002-01-17
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 32
TYPE: PRT
ORGANISM: Pseudomonas sp.
FEATURE:
OTHER INFORMATION: BHY-1(KCTC 0688BP)
US-10-118-047-2

Query Match          83.3%; Score 40; DB 4; Length 32;
Best Local Similarity 77.8%; Pred. No. 2.8;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFQDPOER 9
Db 20 ALFQDPOER 28

RESULT 31
US-10-118-047-2
Sequence 2, Application US/10118047
Publication No. US20040161832A9
GENERAL INFORMATION:
APPLICANT: CHUNG, BONG HYUN
APPLICANT: LEE, EUN GYO
APPLICANT: HAHM, MOON SUN
APPLICANT: RYU, YEON WOO
APPLICANT: LEE, HAN SEUNG
TITLE OF INVENTION: ESTERASE, ITS DNA, ITS OVEREXPRESSION AND PRODUCTION OF
FILE REFERENCE: 05823.0218-00000
CURRENT APPLICATION NUMBER: US/10/118,047
```

```

; CURRENT FILING DATE: 2002-07-03
; PRIOR APPLICATION NUMBER: KR 2002-2809
; PRIOR FILING DATE: 2002-01-17
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: Patent In Ver. 2.1
; SEQ ID NO 2
; LENGTH: 32
; TYPE: PRT
; ORGANISM: Pseudomonas sp.
; FEATURE:
; OTHER INFORMATION: BHY-1 (KCTC 0688BP)
US-10-118-047-2
```

```

Query Match      83.3%; Score 40; DB 4; Length 32;
Best Local Similarity 77.8%; Pred. No. 2.8;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy      1 AMFDDPOER 9
        ||:|||||
Db      20 ALFDDPOER 28
```

```

RESULT 32
US-11-021-949-26
; Sequence 26, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SAMIENTO, CHAMORRO SONOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 26
; LENGTH: 151
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-26
```

```

Query Match      83.3%; Score 40; DB 6; Length 151;
Best Local Similarity 87.5%; Pred. No. 14;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      2 MFQDDPOER 9
        |||||:|
Db      1 MFQDDPOER 8
```

```

RESULT 33
US-10-859-572-2
; Sequence 2, Application US/10859572
; Publication No. US20040219594A1
; GENERAL INFORMATION:
; APPLICANT: Chung, Bong H
; APPLICANT: Lee, Eun G
; APPLICANT: Hahn, Moon S
; APPLICANT: Ryu, Yeon W
; APPLICANT: Lee, Han S
; TITLE OF INVENTION: Esterase, Its DNA, Its Overexpression and Production of
; FILE REFERENCE: 05823.0218-01000
; CURRENT APPLICATION NUMBER: US/10/859,572
; PRIOR FILING DATE: 2004-06-03
; PRIOR APPLICATION NUMBER: 10/118,047
; PRIOR FILING DATE: 2002-04-09
```

```

; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: Patent In Ver. 2.1
; SEQ ID NO 2
; LENGTH: 381
; TYPE: PRT
; ORGANISM: Pseudomonas sp. BHY-1 (KCTC 0688BP)
US-10-859-572-2
```

```

Query Match      83.3%; Score 40; DB 5; Length 381;
Best Local Similarity 77.8%; Pred. No. 36;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy      1 AMFDDPOER 9
        ||:|||||
Db      20 ALFDDPOER 28
```

```

RESULT 34
US-10-751-845-71
; Sequence 71, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; PRIOR FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 71
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-71
```

```

Query Match      81.2%; Score 39; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      3 FQDDPOER 9
        |||||:|
Db      1 FQDDPOER 7
```

```

RESULT 35
US-10-751-845-74
; Sequence 74, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; PRIOR FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 74
```

LENGTH: 10
TYPE: PRT
ORGANISM: Human papilloma virus
US-10-751-845-74

Query Match 81.2%; Score 39; DB 5; Length 10;
Best Local Similarity 100.0%; Pred. No. 1.3;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
1 FODPOER 7

RESULT 36
US-11-021-949-1
Sequence 1, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 1
LENGTH: 25
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-1

Query Match 81.2%; Score 39; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 3.3;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
1 FODPOER 7

RESULT 37
US-10-739-930-9788
Sequence 9788, Application US/10739930
Publication No. US20040216190A1
GENERAL INFORMATION:
APPLICANT: Kovalic, David K.
TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
FILE REFERENCE: 38-21(53377)B
CURRENT APPLICATION NUMBER: US/10/739,930
CURRENT FILING DATE: 2003-12-18
NUMBER OF SEQ ID NOS: 11088
SEQ ID NO 9788
LENGTH: 405
TYPE: PRT
ORGANISM: Triticum aestivum
FEATURE:
OTHER INFORMATION: Clone ID: TRIAE-23APR03-C17806_1.p
US-10-739-930-9788

Query Match 81.2%; Score 39; DB 5; Length 405;
Best Local Similarity 100.0%; Pred. No. 59;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9

Db 275 FODPOER 281

RESULT 38
US-11-021-949-14
Sequence 14, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 14
LENGTH: 149
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-14

Query Match 79.2%; Score 38; DB 6; Length 149;
Best Local Similarity 87.5%; Pred. No. 32;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
1 MFODPOER 8

RESULT 39
US-11-021-949-27
Sequence 27, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 27
LENGTH: 150
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-27

Query Match 77.1%; Score 37; DB 6; Length 150;
Best Local Similarity 75.0%; Pred. No. 50;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
1 MFODPOER 8

RESULT 40

```
US-10-159-257A-149
; Sequence 149, Application US/10159257A
; Publication No. US20040161828A1
; GENERAL INFORMATION:
; APPLICANT: LIU, MEN
; APPLICANT: SHEN, BEN
; APPLICANT: CHRISTENSON, STEVEN D.
; APPLICANT: STANDAGE, SCOTT
; TITLE OF INVENTION: GENE CLUSTER FOR PRODUCTION OF THE ENEDIYNE ANTITUMOR
; FILE REFERENCE: 4077-896020US
; CURRENT APPLICATION NUMBER: US/10/159,257A
; PRIOR FILING DATE: 2002-05-31
; PRIOR APPLICATION NUMBER: 09/478,188
; PRIOR FILING DATE: 2000-01-05
; PRIOR APPLICATION NUMBER: 60/115,434
; PRIOR FILING DATE: 1999-01-06
; NUMBER OF SEQ ID NOS: 207
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 149
; LENGTH: 521
; TYPE: PRT
; ORGANISM: Streptomyces globisporus
; US-10-159-257A-149

Query Match      77.1%; Score 37; DB 4; Length 521;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 MFQDPOER 9
      |||:|:|
Db      133 MFEDPKER 140

RESULT 41
US-10-159-257A-181
; Sequence 181, Application US/10159257A
; Publication No. US20040161828A1
; GENERAL INFORMATION:
; APPLICANT: SHEN, BEN
; APPLICANT: LIU, MEN
; APPLICANT: CHRISTENSON, STEVEN D.
; APPLICANT: STANDAGE, SCOTT
; TITLE OF INVENTION: GENE CLUSTER FOR PRODUCTION OF THE ENEDIYNE ANTITUMOR
; FILE REFERENCE: 4077-896020US
; CURRENT APPLICATION NUMBER: US/10/159,257A
; CURRENT FILING DATE: 2002-05-31
; PRIOR FILING DATE: 09/478,188
; PRIOR FILING DATE: 2000-01-05
; PRIOR APPLICATION NUMBER: 60/115,434
; PRIOR FILING DATE: 1999-01-06
; NUMBER OF SEQ ID NOS: 207
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 181
; LENGTH: 521
; TYPE: PRT
; ORGANISM: Streptomyces globisporus
; FEATURE:
; OTHER INFORMATION: orf2
; US-10-159-257A-181

Query Match      77.1%; Score 37; DB 4; Length 521;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 MFQDPOER 9
      |||:|:|
Db      133 MFEDPKER 140

RESULT 42
US-10-425-114-68828

; Sequence 68828, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: LIU, Jindong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; FILE REFERENCE: 38-21(5313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 68828
; LENGTH: 354
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-ZMFLMO17223F12_F11.pep
; US-10-425-114-68828

Query Match      75.0%; Score 36; DB 4; Length 354;
Best Local Similarity 85.7%; Pred. No. 1.9e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 FQDPQER 9
      |||||:|
Db      228 FQDPQER 234

RESULT 43
US-10-425-115-330168
; Sequence 330168, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; FILE REFERENCE: 38-21(5322)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 330168
; LENGTH: 354
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_64213C.1.pep
; US-10-425-115-330168

Query Match      75.0%; Score 36; DB 4; Length 354;
Best Local Similarity 85.7%; Pred. No. 1.9e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 FQDPQER 9
      |||||:|
Db      228 FQDPQER 234

RESULT 44
US-10-425-115-341903
; Sequence 341903, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
```



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; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 341903
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Zea mays
; PEATRE:
; OTHER INFORMATION: Clone ID: MTT4577_74983C.1.pep
; US-10-425-115-341903

Query Match          72.9%; Score 35; DB 4; Length 129;
Best Local Similarity 75.0%; Pred. No. 16+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      2 MFQDPOER 9
      : |||||
      : 41 MFSDPOKR 48

Db

RESULT 45
US-10-282-122A-68570
; Sequence 68570, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Karl
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELITRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20
; PRIOR APPLICATION NUMBER: 60/191,078
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: 60/206,848
; PRIOR FILING DATE: 2000-05-23
; PRIOR APPLICATION NUMBER: 60/207,727
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: 60/230,335
; PRIOR FILING DATE: 2000-09-06
; PRIOR APPLICATION NUMBER: 60/230,347
; PRIOR FILING DATE: 2000-09-09
; PRIOR APPLICATION NUMBER: 60/242,578
; PRIOR FILING DATE: 2000-10-23
; PRIOR APPLICATION NUMBER: 60/253,625
; PRIOR FILING DATE: 2000-11-27
; PRIOR APPLICATION NUMBER: 60/257,931
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: 60/267,636
; PRIOR FILING DATE: 2001-02-09
; PRIOR APPLICATION NUMBER: 60/269,308
; PRIOR FILING DATE: 2001-02-16
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 78614
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 68570
; LENGTH: 1144
; TYPE: PRT
; ORGANISM: Proteus mirabilis
; US-10-282-122A-68570
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Query Match          72.9%; Score 35; DB 4; Length 1144;
Best Local Similarity 75.0%; Pred. No. 9.8e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      2 MFQDPOER 9
      : |||||
      : 291 LFQDPOER 298

Db

RESULT 46
US-09-864-761-45512
; Sequence 45512, Application US/09864761
; Patent No. US20020048763A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharon G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; APPLICANT: Chen, Wensheng
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR
; TITLE OF INVENTION: GENE EXPRESSION ANALYSIS BY MICROARRAY
; FILE REFERENCE: Aeomica-X-1
; CURRENT APPLICATION NUMBER: US/09/864,761
; CURRENT FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: US 60/180,312
; PRIOR FILING DATE: 2000-02-04
; PRIOR APPLICATION NUMBER: US 60/207,456
; PRIOR FILING DATE: 2000-05-26
; PRIOR APPLICATION NUMBER: US 09/632,366
; PRIOR FILING DATE: 2000-08-03
; PRIOR APPLICATION NUMBER: GB 24263.6
; PRIOR FILING DATE: 2000-10-04
; PRIOR APPLICATION NUMBER: US 60/236,359
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: PCT/US01/00666
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00667
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00664
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00669
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00665
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00668
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00663
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00662
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00661
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: PCT/US01/00670
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: US 60/234,687
; PRIOR FILING DATE: 2000-09-21
; PRIOR APPLICATION NUMBER: US 09/608,408
; PRIOR FILING DATE: 2000-06-30
; PRIOR APPLICATION NUMBER: US 09/774,203
; PRIOR FILING DATE: 2001-01-29
; NUMBER OF SEQ ID NOS: 49117
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 45512
; LENGTH: 29
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO AC010363.3
; OTHER INFORMATION: EXPRESSED IN PLACENTA, SIGNAL = 0.71
; OTHER INFORMATION: EXPRESSED IN FETAL LIVER, SIGNAL = 0.69
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.76
; OTHER INFORMATION: EXPRESSED IN BONE MARROW, SIGNAL = 0.84
; OTHER INFORMATION: EXPRESSED IN LUNG, SIGNAL = 0.87
; OTHER INFORMATION: EXPRESSED IN BRAIN, SIGNAL = 0.77
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US-09-864-761-45512

Query Match 70.8%; Score 34; DB 3; Length 29;
Best Local Similarity 55.6%; Pred. No. 34;
Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOQ 9
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DB 13 AVFODPODK 21

RESULT 47

US-10-504-582-119
; Sequence 119, Application US/10504582
; Publication No. US20050176943A1
; GENERAL INFORMATION:
; APPLICANT: Yoshitake NISHIMUNE
; APPLICANT: Hiromitsu TANAKA
; APPLICANT: Masami NOZAKI
; TITLE OF INVENTION: Mouse spermatogenesis genes, mutations of male infertility-relate
; FILE REFERENCE: 2004-1256A/WMC/00653
; CURRENT FILING DATE: 2004-08-13
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: JP2002-36649
; PRIOR FILING DATE: 2002-12-27
; NUMBER OF SEQ ID NOS: 183
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO: 119
; LENGTH: 137
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-504-582-119

Query Match 70.8%; Score 34; DB 5; Length 137;
Best Local Similarity 85.7%; Pred. No. 1.7e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOQ 7
|:|||||
DB 44 AVFODPO 50

RESULT 48

US-10-425-115-203539
; Sequence 203539, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovacic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO: 203539
; LENGTH: 204
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; LOCATION: (1)-(204)
; OTHER INFORMATION: unsure at all Xaa locations
; OTHER INFORMATION: Clone ID: MRT4577_117217C.1.psp
US-10-425-115-203539

Query Match 70.8%; Score 34; DB 4; Length 204;

Best Local Similarity 75.0%; Pred. No. 2.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
||:||||
DB 165 MFODPOER 172

RESULT 49

US-09-815-453-8
; Sequence 8, Application US/09815453
; Publication No. US20030166519A1
; GENERAL INFORMATION:
; APPLICANT: Huynh, The Hung
; TITLE OF INVENTION: and Uses therefor
; FILE REFERENCE: 32061000000
; CURRENT FILING DATE: 2001-03-22
; PRIOR FILING DATE: 2000-08-15
; PRIOR APPLICATION NUMBER: AU PQ9470
; PRIOR FILING DATE: 2000-05-25
; PRIOR APPLICATION NUMBER: AU PQ7732
; PRIOR FILING DATE: 2000-05-24
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 8
; LENGTH: 456
; TYPE: PRT
; ORGANISM: Mouse
US-09-815-453-8

Query Match 70.8%; Score 34; DB 3; Length 456;
Best Local Similarity 85.7%; Pred. No. 5.8e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPO 7
|:|||||
DB 354 AVFODPO 360

RESULT 50

US-10-087-192-867
; Sequence 867, Application US/10087192
; Publication No. US20020182586A1
; GENERAL INFORMATION:
; APPLICANT: Morris, David W.
; APPLICANT: Engelhard, Eric K.
; TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR
; FILE REFERENCE: 529452000122
; CURRENT FILING DATE: 2002-03-01
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: US 09/798,586
; PRIOR FILING DATE: 2001-03-02
; NUMBER OF SEQ ID NOS: 2059
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 867
; LENGTH: 484
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-087-192-867

Query Match 70.8%; Score 34; DB 4; Length 484;
Best Local Similarity 85.7%; Pred. No. 6.2e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPO 7
|:|||||
DB 382 AVFODPO 388

Tue May 9 09:28:04 2006

us-08-170-344-1.rapbm

Page 21

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OM protein - protein search, using sw model

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(without alignments)
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Perfect score: 48
Sequence: 1 AMFDPQER 9

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Gapop 10.0 , Gapext 0.5

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

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2: /SIDSS/ptodata/1/pubppaa/US06_NEW_PUB.pep.*
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5: /SIDSS/ptodata/1/pubppaa/PCR_NEW_PUB.pep.*
6: /SIDSS/ptodata/1/pubppaa/US09_NEW_PUB.pep.*
7: /SIDSS/ptodata/1/pubppaa/US10_NEW_PUB.pep.*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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2	48	100.0	11	9	US-10-530-061-482
3	48	100.0	15	9	US-10-530-061-1702
4	48	100.0	158	11	US-11-206-138-3
5	48	100.0	256	11	US-11-192-923A-2
6	44	91.7	11	9	US-10-530-061-483
7	44	91.7	151	9	US-10-530-253-13
8	44	91.7	248	9	US-10-530-253-1
9	44	91.7	248	9	US-10-530-253-3
10	44	91.7	248	9	US-10-530-253-5
11	39	81.2	248	9	US-10-530-253-7
12	39	81.2	248	9	US-10-530-253-9
13	39	81.2	248	9	US-10-530-253-11
14	38	79.2	149	9	US-10-530-253-18
15	36	75.0	567	11	US-11-079-463-7932
16	32	66.7	144	11	US-11-096-568A-18478
17	32	66.7	158	9	US-10-530-253-26
18	32	66.7	167	11	US-11-096-568A-18476
19	32	66.7	288	11	US-11-188-298-16639
20	32	66.7	325	8	US-10-505-928-327
21	32	66.7	325	10	US-11-301-554-1816

22	66.7	329	11	US-11-188-298-13197	Sequence 13197, A
23	66.7	399	9	US-10-821-234-1163	Sequence 1163, Ap
24	66.7	1341	11	US-11-096-568A-32147	Sequence 32147, A
25	66.7	1372	11	US-11-096-568A-32146	Sequence 32146, A
26	66.7	1393	11	US-11-096-568A-32145	Sequence 32145, A
27	64.6	152	9	US-10-530-253-39	Sequence 39, Appl
28	64.6	445	11	US-11-096-568A-18941	Sequence 18941, A
29	64.6	447	9	US-10-491-468-5	Sequence 5, Appl1
30	64.6	470	11	US-11-096-568A-18940	Sequence 18940, A
31	64.6	486	11	US-11-096-568A-18939	Sequence 18939, A
32	62.5	149	9	US-10-530-253-17	Sequence 17, Appl
33	62.5	149	9	US-10-530-253-24	Sequence 24, Appl
34	62.5	158	9	US-10-530-253-20	Sequence 20, Appl
35	62.5	160	9	US-10-499-253-25	Sequence 25, Appl
36	62.5	293	9	US-10-499-246-17	Sequence 18, Appl
37	62.5	301	9	US-10-499-246-18	Sequence 1478, Ap
38	62.5	309	9	US-10-506-454-1478	Sequence 1328, Ap
39	62.5	326	9	US-10-793-626-3338	Sequence 132, Ap
40	62.5	365	11	US-11-082-389-132	Sequence 29694, A
41	62.5	367	11	US-11-096-568A-29694	Sequence 29693, A
42	62.5	380	11	US-11-096-568A-29693	Sequence 29692, A
43	62.5	416	11	US-11-096-568A-29692	Sequence 130, Ap
44	62.5	479	11	US-11-082-389-130	Sequence 16827, A
45	62.5	510	11	US-11-188-298-16827	Sequence 6591, Ap
46	62.5	533	11	US-11-079-463-6591	Sequence 20601, A
47	62.5	543	11	US-11-188-298-20601	Sequence 22376, A
48	62.5	548	11	US-11-188-298-22276	Sequence 2436, Ap
49	62.5	601	11	US-11-087-039-2496	Sequence 11892, A
50	62.5	648	11	US-11-087-039-11892	Sequence 648, Appl
51	62.5	762	9	US-10-506-454-648	Sequence 34, Appl
52	62.5	966	9	US-10-532-482-34	Sequence 34, Appl
53	62.5	979	9	US-10-532-482-37	Sequence 50, Appl
54	62.5	982	9	US-10-532-482-55	Sequence 60, Appl
55	62.5	985	9	US-10-532-482-60	Sequence 45, Appl
56	62.5	987	9	US-10-532-482-45	Sequence 45, Appl
57	62.5	993	9	US-10-532-482-39	Sequence 51, Appl
58	62.5	993	9	US-10-532-482-51	Sequence 51, Appl
59	62.5	995	9	US-10-532-482-47	Sequence 58, Appl
60	62.5	997	9	US-10-532-482-58	Sequence 43, Appl
61	62.5	1001	9	US-10-532-482-43	Sequence 41, Appl
62	62.5	1004	9	US-10-532-482-41	Sequence 59, Appl
63	62.5	1006	9	US-10-532-482-59	Sequence 54, Appl
64	62.5	1006	9	US-10-532-482-54	Sequence 53, Appl
65	62.5	1006	9	US-10-532-482-53	Sequence 31, Appl
66	62.5	1009	9	US-10-532-482-31	Sequence 32, Appl
67	62.5	1013	9	US-10-532-482-32	Sequence 50, Appl
68	62.5	1016	9	US-10-532-482-30	Sequence 50, Appl
69	62.5	1016	9	US-10-532-482-57	Sequence 50, Appl
70	62.5	1017	9	US-10-532-482-30	Sequence 50, Appl
71	62.5	1020	9	US-10-532-482-35	Sequence 35, Appl
72	62.5	1029	9	US-10-532-482-52	Sequence 49, Appl
73	62.5	1032	9	US-10-532-482-44	Sequence 56, Appl
74	62.5	1033	9	US-10-532-482-56	Sequence 56, Appl
75	62.5	1033	9	US-10-532-482-36	Sequence 44, Appl
76	62.5	1041	9	US-10-532-482-44	Sequence 38, Appl
77	62.5	1047	9	US-10-532-482-38	Sequence 46, Appl
78	62.5	1049	9	US-10-532-482-46	Sequence 42, Appl
79	62.5	1055	9	US-10-532-482-42	Sequence 40, Appl
80	62.5	1057	9	US-10-532-482-40	Sequence 18, Appl
81	62.5	1061	11	US-11-059-814-18	Sequence 48, Appl
82	62.5	1071	9	US-10-532-482-18	Sequence 97, Appl
83	62.5	1176	9	US-10-821-234-897	Sequence 22, Appl
84	60.4	148	9	US-10-530-253-12	Sequence 16, Appl
85	60.4	149	9	US-10-530-253-16	Sequence 21, Appl
86	60.4	151	9	US-10-530-253-21	Sequence 23, Appl
87	60.4	155	9	US-10-530-253-23	Sequence 1734, Ap
88	60.4	191	11	US-11-172-740-1134	Sequence 1735, Ap
89	60.4	191	11	US-11-172-740-1135	Sequence 1740, Ap
90	60.4	192	11	US-11-172-740-1140	Sequence 1741, Ap
91	60.4	192	11	US-11-172-740-1141	Sequence 1742, Ap
92	60.4	192	11	US-11-172-740-1142	Sequence 1743, Ap
93	60.4	192	11	US-11-172-740-1143	Sequence 1744, Ap
94	60.4	192	11	US-11-172-740-1144	Sequence 1744, Ap

95	29	60.4	220	9	US-10-506-454-1038	Sequence 1038, Ap
96	29	60.4	229	11	US-11-072-512-3781	Sequence 3781, Ap
97	29	60.4	252	9	US-10-506-454-764	Sequence 764, App
98	29	60.4	278	11	US-11-165-067A-5	Sequence 5, Appl1
99	29	60.4	319	9	US-10-467-657-3252	Sequence 3252, Ap
100	29	60.4	336	11	US-11-188-298-12437	Sequence 12437, A
101	29	60.4	352	11	US-11-188-298-6720	Sequence 6720, Ap
102	29	60.4	369	11	US-11-096-568A-11196	Sequence 11196, A
103	29	60.4	397	11	US-11-072-512-2919	Sequence 2919, Ap
104	29	60.4	460	11	US-11-096-568A-24725	Sequence 24725, A
105	29	60.4	488	11	US-11-079-463-6937	Sequence 6937, Ap
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107	29	60.4	501	11	US-11-188-298-7723	Sequence 7723, Ap
108	29	60.4	512	9	US-10-821-234-1032	Sequence 1032, Ap
109	29	60.4	520	11	US-11-087-099-11701	Sequence 11701, A
110	29	60.4	520	11	US-11-188-298-21871	Sequence 21871, A
111	29	60.4	530	11	US-11-096-568A-24724	Sequence 24724, A
112	29	60.4	536	11	US-11-096-568A-24773	Sequence 24773, A
113	29	60.4	547	9	US-10-242-586-40	Sequence 40, Appl
114	29	60.4	547	9	US-10-242-902-40	Sequence 40, Appl
115	29	60.4	547	9	US-10-243-116-40	Sequence 40, Appl
116	29	60.4	547	9	US-10-243-136-40	Sequence 40, Appl
117	29	60.4	547	9	US-10-243-189-40	Sequence 40, Appl
118	29	60.4	547	9	US-10-243-215-40	Sequence 40, Appl
119	29	60.4	547	9	US-10-243-236-40	Sequence 40, Appl
120	29	60.4	547	9	US-10-243-298-40	Sequence 40, Appl
121	29	60.4	547	9	US-10-243-304-40	Sequence 40, Appl
122	29	60.4	547	9	US-10-243-338-40	Sequence 40, Appl
123	29	60.4	547	9	US-10-243-345-40	Sequence 40, Appl
124	29	60.4	547	9	US-10-243-357-40	Sequence 40, Appl
125	29	60.4	547	9	US-10-245-083-40	Sequence 40, Appl
126	29	60.4	547	9	US-10-247-015-40	Sequence 40, Appl
127	29	60.4	577	11	US-11-087-099-5216	Sequence 5216, Ap
128	29	60.4	577	11	US-11-188-298-4767	Sequence 4767, Ap
129	29	60.4	611	11	US-11-096-568A-9419	Sequence 9419, Ap
130	29	60.4	684	11	US-11-096-568A-9418	Sequence 9418, Ap
131	29	60.4	711	11	US-11-096-568A-9417	Sequence 9417, Ap
132	29	60.4	1076	9	US-10-467-657-7916	Sequence 7916, Ap
133	29	60.4	1199	9	US-10-204-639-32	Sequence 32, Appl
134	29	60.4	3073	11	US-11-143-980-50	Sequence 50, Appl
135	29	60.4	5712	11	US-11-143-980-47	Sequence 47, Appl
136	29	60.4	7968	11	US-11-143-980-49	Sequence 49, Appl
137	28	58.3	83	11	US-11-079-463-9743	Sequence 9743, Ap
138	28	58.3	117	11	US-11-172-740-810	Sequence 810, App
139	28	58.3	151	11	US-11-082-389-108	Sequence 108, App
140	28	58.3	158	9	US-10-530-253-15	Sequence 15, Appl
141	28	58.3	162	11	US-11-134-811-10	Sequence 10, Appl
142	28	58.3	223	11	US-11-079-463-8627	Sequence 8627, Ap
143	28	58.3	246	11	US-11-096-568A-23753	Sequence 23753, A
144	28	58.3	268	11	US-11-046-004-437	Sequence 43, Appl
145	28	58.3	269	11	US-11-188-298-857	Sequence 857, App
146	28	58.3	273	11	US-11-096-568A-11741	Sequence 11741, A
147	28	58.3	277	11	US-11-072-512-2535	Sequence 2535, Ap
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153	28	58.3	304	11	US-11-096-568A-29450	Sequence 29450, A
154	28	58.3	304	11	US-11-079-463-6327	Sequence 6327, Ap
155	28	58.3	305	11	US-11-096-568A-21133	Sequence 2133, Ap
156	28	58.3	307	11	US-11-188-298-5745	Sequence 5745, Ap
157	28	58.3	315	11	US-11-096-568A-11831	Sequence 11831, A
158	28	58.3	320	11	US-11-096-568A-1218	Sequence 1218, Ap
159	28	58.3	342	11	US-11-096-568A-29449	Sequence 29449, A
160	28	58.3	348	11	US-11-096-568A-22775	Sequence 22775, A
161	28	58.3	350	11	US-11-096-568A-22774	Sequence 22774, A
162	28	58.3	355	11	US-11-108-528-18	Sequence 18, Appl
163	28	58.3	358	11	US-11-087-099-2705	Sequence 2705, Ap
164	28	58.3	359	11	US-11-087-099-9152	Sequence 9152, Ap
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166	28	58.3	360	9	US-10-195-883-28	Sequence 28, Appl
167	28	58.3	360	9	US-10-195-886-28	Sequence 28, Appl
168	28	58.3	360	9	US-10-195-889-28	Sequence 28, Appl
169	28	58.3	360	11	US-11-188-298-1256	Sequence 3256, Ap
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171	28	58.3	362	11	US-11-096-568A-11739	Sequence 11739, A
172	28	58.3	364	11	US-11-087-099-9542	Sequence 9542, A
173	28	58.3	366	11	US-11-087-099-697	Sequence 697, App
174	28	58.3	366	11	US-11-087-099-6470	Sequence 6470, A
175	28	58.3	366	11	US-11-096-568A-22773	Sequence 22773, A
176	28	58.3	367	11	US-11-087-099-6836	Sequence 6836, Ap
177	28	58.3	370	11	US-11-087-099-6834	Sequence 6834, Ap
178	28	58.3	371	11	US-11-087-099-6732	Sequence 6732, Ap
179	28	58.3	371	11	US-11-087-099-6730	Sequence 6730, Ap
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181	28	58.3	386	11	US-11-098-686-10223	Sequence 10223, A
182	28	58.3	392	11	US-11-096-568A-12127	Sequence 12127, Ap
183	28	58.3	405	9	US-10-506-454-650	Sequence 650, App
184	28	58.3	406	11	US-11-075-185-23	Sequence 23, Appl
185	28	58.3	437	9	US-10-467-657-4088	Sequence 4088, Ap
186	28	58.3	439	11	US-11-096-568A-1216	Sequence 1216, Ap
187	28	58.3	445	9	US-10-821-234-1606	Sequence 1606, Ap
188	28	58.3	463	11	US-11-024-959-507	Sequence 507, App
189	28	58.3	517	11	US-11-087-099-11344	Sequence 11344, A
190	28	58.3	517	11	US-11-188-298-10464	Sequence 10464, A
191	28	58.3	533	11	US-11-087-099-5854	Sequence 5854, Ap
192	28	58.3	548	9	US-10-213-535-24	Sequence 24, Appl
193	28	58.3	562	9	US-10-508-263-24	Sequence 24, Appl
194	28	58.3	599	11	US-11-188-298-16912	Sequence 16912, A
195	28	58.3	630	9	US-10-467-657-7014	Sequence 7014, Ap
196	28	58.3	879	11	US-11-096-568A-30815	Sequence 30815, A
197	28	58.3	929	9	US-10-467-657-5656	Sequence 5656, Ap
198	28	58.3	936	11	US-11-072-512-2621	Sequence 2621, Ap
199	28	58.3	964	11	US-11-103-957-13	Sequence 13, Appl
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202	28	58.3	2197	11	US-11-075-185-8	Sequence 8, Appl1
203	28	58.3	2238	9	US-10-330-773-32	Sequence 42, Appl
204	28	58.3	2910	9	US-10-330-773-39	Sequence 39, Appl
205	28	58.3	3689	11	US-11-075-185-4	Sequence 4, Appl1
206	28	58.3	7102	11	US-11-143-980-48	Sequence 48, Appl
207	27	56.2	82	9	US-10-506-454-962	Sequence 962, App
208	27	56.2	84	9	US-10-467-657-5404	Sequence 5404, Ap
209	27	56.2	98	11	US-11-087-099-4711	Sequence 4711, Ap
210	27	56.2	147	9	US-10-991-285-895	Sequence 895, App
211	27	56.2	157	9	US-10-506-454-1586	Sequence 1586, Ap
212	27	56.2	170	11	US-11-232-406A-26	Sequence 26, Appl
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214	27	56.2	184	11	US-11-096-568A-25177	Sequence 25177, A
215	27	56.2	184	11	US-11-153-071-10	Sequence 10, Appl
216	27	56.2	185	11	US-11-087-099-1152	Sequence 1152, Ap
217	27	56.2	190	11	US-11-096-568A-3909	Sequence 3909, Ap
218	27	56.2	197	11	US-11-087-099-8247	Sequence 8247, Ap
219	27	56.2	197	9	US-10-455-772-108	Sequence 108, App
220	27	56.2	200	11	US-11-087-099-12079	Sequence 12079, A
221	27	56.2	201	11	US-11-082-389-130	Sequence 138, App
222	27	56.2	205	11	US-11-045-004-398	Sequence 398, App
223	27	56.2	213	11	US-11-096-568A-16401	Sequence 16401, A
224	27	56.2	213	11	US-11-096-568A-28213	Sequence 28213, A
225	27	56.2	219	11	US-11-096-568A-12944	Sequence 12944, A
226	27	56.2	223	11	US-11-096-568A-16400	Sequence 16400, A
227	27	56.2	235	11	US-11-096-568A-12943	Sequence 12943, A
228	27	56.2	242	9	US-10-467-657-438	Sequence 438, App
229	27	56.2	245	9	US-10-506-454-1149	Sequence 1149, Ap
230	27	56.2	248	11	US-11-096-568A-3390	Sequence 3390, Ap
231	27	56.2	254	9	US-10-506-454-1885	Sequence 1685, Ap
232	27	56.2	256	11	US-11-096-568A-3389	Sequence 3389, Ap
233	27	56.2	259	11	US-11-096-568A-21634	Sequence 21634, A
234	27	56.2	260	11	US-11-096-568A-28212	Sequence 28212, A
235	27	56.2	261	9	US-10-617-034A-12	Sequence 12, Appl
236	27	56.2	271	11	US-11-045-004-3691	Sequence 2691, Ap
237	27	56.2	284	9	US-10-131-828A-118	Sequence 118, App
238	27	56.2	284	9	US-10-973-115B-118	Sequence 118, App
239	27	56.2	284	9	US-10-137-873A-118	Sequence 118, App
240	27	56.2	284	9	US-10-152-370-118	Sequence 118, App

241	27	56.2	284	11	US-11-087-099-11149	Sequence 11149, A	314	27	56.2	566	11	US-11-072-512-2407	Sequence 2407, Ap
242	27	56.2	284	11	US-11-230-153-118	Sequence 118, App	315	27	56.2	577	11	US-11-082-389-134	Sequence 134, App
243	27	56.2	286	11	US-11-089-551A-27	Sequence 27, Appl	316	27	56.2	588	9	US-10-784-004-459	Sequence 459, App
244	27	56.2	288	11	US-11-096-568A-3908	Sequence 3908, Ap	317	27	56.2	593	9	US-10-784-004-777	Sequence 777, App
245	27	56.2	289	11	US-11-096-568A-3907	Sequence 3907, Ap	318	27	56.2	599	9	US-10-455-772-110	Sequence 110, App
246	27	56.2	297	11	US-11-096-568A-12942	Sequence 12942, A	319	27	56.2	605	9	US-10-455-772-106	Sequence 106, App
247	27	56.2	297	11	US-11-087-099-3875	Sequence 3875, Ap	320	27	56.2	613	9	US-10-455-772-112	Sequence 112, App
248	27	56.2	316	11	US-11-188-298-513	Sequence 513, App	321	27	56.2	637	11	US-11-113-837-4	Sequence 4, Appl
249	27	56.2	317	11	US-11-087-099-7583	Sequence 7583, Ap	322	27	56.2	638	9	US-10-455-772-104	Sequence 104, App
250	27	56.2	335	11	US-11-188-298-9304	Sequence 9304, Ap	323	27	56.2	672	11	US-11-079-463-9531	Sequence 9531, Ap
251	27	56.2	348	11	US-11-188-298-14107	Sequence 14107, A	324	27	56.2	691	11	US-11-188-298-19528	Sequence 19528, A
252	27	56.2	349	11	US-11-129-143-45	Sequence 45, Appl	325	27	56.2	693	11	US-11-188-298-3925	Sequence 3925, Ap
253	27	56.2	349	11	US-11-096-568A-7422	Sequence 7422, Ap	326	27	56.2	707	9	US-10-784-004-1222	Sequence 1222, Ap
254	27	56.2	353	11	US-11-087-099-10589	Sequence 10589, A	327	27	56.2	719	11	US-11-063-243-33	Sequence 33, Appl
255	27	56.2	354	11	US-11-096-568A-10863	Sequence 10863, A	328	27	56.2	744	11	US-11-052-454A-1112	Sequence 112, App
256	27	56.2	355	11	US-11-087-099-5542	Sequence 5542, Ap	329	27	56.2	849	11	US-11-188-298-7198	Sequence 7198, Ap
257	27	56.2	356	11	US-11-087-099-846	Sequence 846, App	330	27	56.2	979	9	US-10-517-622-2	Sequence 2, Appl
258	27	56.2	356	11	US-11-087-099-4390	Sequence 4390, Ap	331	27	56.2	1056	8	US-10-505-928-225	Sequence 225, App
259	27	56.2	356	11	US-11-087-099-5763	Sequence 5763, Ap	332	27	56.2	1126	11	US-11-075-185-3	Sequence 3, Appl
260	27	56.2	356	11	US-11-087-099-5793	Sequence 5793, Ap	333	27	56.2	1170	8	US-10-511-937-1007	Sequence 3007, Ap
261	27	56.2	356	11	US-11-087-099-7270	Sequence 7270, Ap	334	27	56.2	1170	11	US-11-080-026-2	Sequence 2, Appl
262	27	56.2	356	11	US-11-087-099-7589	Sequence 7589, Ap	335	27	56.2	1170	11	US-11-107-028-4	Sequence 4, Appl
263	27	56.2	356	11	US-11-087-099-11044	Sequence 11044, A	336	27	56.2	1246	11	US-11-288-720-17	Sequence 17, Appl
264	27	56.2	357	11	US-11-087-099-3758	Sequence 3758, Ap	337	27	56.2	1255	9	US-10-770-726-62	Sequence 62, Appl
265	27	56.2	357	11	US-11-087-099-5974	Sequence 5974, Ap	338	27	56.2	1255	9	US-11-022-562-213	Sequence 213, App
266	27	56.2	357	11	US-11-087-099-9311	Sequence 9311, Ap	339	27	56.2	1255	11	US-11-113-302-10	Sequence 10, Appl
267	27	56.2	358	11	US-11-087-099-8981	Sequence 8981, Ap	340	27	56.2	1255	11	US-11-033-039-553	Sequence 53, App
268	27	56.2	359	11	US-11-087-099-7895	Sequence 7895, Ap	341	27	56.2	1255	11	US-11-135-288-9	Sequence 9, Appl
269	27	56.2	360	9	US-10-506-454-47	Sequence 47, Appl	342	27	56.2	1255	11	US-11-202-516-4	Sequence 4, Appl
270	27	56.2	365	9	US-10-770-726-69	Sequence 69, Appl	343	27	56.2	1258	11	US-11-175-405-2	Sequence 2, Appl
271	27	56.2	367	9	US-10-525-674-2	Sequence 2, Appl	344	27	56.2	1358	11	US-11-033-039-930	Sequence 930, App
272	27	56.2	368	11	US-11-188-298-19925	Sequence 19925, A	345	27	56.2	1336	9	US-10-912-971-10	Sequence 10, Appl
273	27	56.2	369	11	US-10-517-939-64	Sequence 64, Appl	346	27	56.2	1483	11	US-11-188-298-10839	Sequence 10839, A
274	27	56.2	384	11	US-11-096-568A-10862	Sequence 10862, A	347	27	56.2	1536	9	US-11-087-099-1921	Sequence 1921, A
275	27	56.2	395	11	US-11-096-568A-22837	Sequence 22837, A	348	27	56.2	9	9	US-10-530-061-375	Sequence 375, App
276	27	56.2	402	11	US-11-087-099-5843	Sequence 5843, Ap	349	26	54.2	10	9	US-10-530-061-187	Sequence 187, App
277	27	56.2	412	11	US-11-096-568A-7421	Sequence 7421, Ap	350	26	54.2	10	9	US-10-530-061-905	Sequence 905, App
278	27	56.2	413	11	US-11-096-568A-7420	Sequence 7420, Ap	351	26	54.2	14	11	US-11-107-539-29	Sequence 29, Appl
279	27	56.2	423	11	US-11-096-568A-29642	Sequence 29642, A	352	26	54.2	19	9	US-10-503-575-258	Sequence 258, App
280	27	56.2	431	8	US-10-505-928-465	Sequence 465, App	353	26	54.2	48	9	US-10-948-571-63	Sequence 63, Appl
281	27	56.2	431	9	US-10-995-561-807	Sequence 807, App	354	26	54.2	69	11	US-11-045-004-3303	Sequence 2303, Ap
282	27	56.2	431	9	US-10-995-561-808	Sequence 808, App	355	26	54.2	84	11	US-11-124-607-8	Sequence 8, Appl
283	27	56.2	431	9	US-10-504-287-2	Sequence 2, Appl	356	26	54.2	90	9	US-10-467-657-3624	Sequence 3624, Ap
284	27	56.2	431	10	US-11-183-218-34	Sequence 34, Appl	357	26	54.2	104	11	US-11-096-568A-17427	Sequence 17427, A
285	27	56.2	431	11	US-11-186-284-161	Sequence 161, App	358	26	54.2	110	11	US-11-096-568A-11974	Sequence 11974, A
286	27	56.2	431	11	US-11-183-205-34	Sequence 34, Appl	359	26	54.2	149	11	US-11-096-568A-11573	Sequence 11573, A
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288	27	56.2	448	9	US-10-763-712A-112	Sequence 112, App	361	26	54.2	158	9	US-10-703-799B-774	Sequence 774, App
289	27	56.2	449	9	US-10-506-454-1070	Sequence 1070, Ap	362	26	54.2	158	11	US-11-082-389-356	Sequence 356, App
290	27	56.2	450	9	US-10-506-454-1537	Sequence 1537, Ap	363	26	54.2	160	9	US-10-467-657-4072	Sequence 4072, Ap
291	27	56.2	452	11	US-11-087-099-588	Sequence 588, App	364	26	54.2	166	11	US-11-188-298-3373	Sequence 3373, Ap
292	27	56.2	452	11	US-11-087-099-12236	Sequence 12236, A	365	26	54.2	167	11	US-11-229-240-2	Sequence 2, Appl
293	27	56.2	452	11	US-11-188-298-638	Sequence 638, App	366	26	54.2	169	11	US-11-096-568A-15465	Sequence 26565, A
294	27	56.2	452	11	US-11-188-298-22347	Sequence 22347, A	367	26	54.2	171	11	US-11-096-568A-16441	Sequence 16441, A
295	27	56.2	457	9	US-10-995-561-797	Sequence 797, App	368	26	54.2	180	11	US-11-264-096-2116	Sequence 2116, Ap
296	27	56.2	462	11	US-11-096-568A-22836	Sequence 22836, A	369	26	54.2	181	11	US-11-264-728-26	Sequence 26, Appl
297	27	56.2	467	9	US-10-519-238-2	Sequence 2, Appl	370	26	54.2	182	11	US-11-096-568A-3775	Sequence 3775, Ap
298	27	56.2	469	11	US-11-087-099-11321	Sequence 11321, A	371	26	54.2	183	11	US-11-096-568A-3774	Sequence 3774, Ap
299	27	56.2	476	11	US-11-031-206-201	Sequence 201, App	372	26	54.2	183	11	US-11-188-298-7633	Sequence 7633, Ap
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301	27	56.2	480	11	US-11-188-298-12730	Sequence 12730, A	374	26	54.2	191	11	US-11-172-740-1736	Sequence 1736, Ap
302	27	56.2	481	11	US-11-096-568A-22835	Sequence 22835, A	375	26	54.2	192	11	US-11-172-740-1739	Sequence 1739, Ap
303	27	56.2	484	11	US-11-087-099-2542	Sequence 2542, Ap	376	26	54.2	196	11	US-11-079-463-8368	Sequence 8368, Ap
304	27	56.2	484	11	US-11-188-298-2423	Sequence 2423, Ap	377	26	54.2	199	11	US-11-043-788-378	Sequence 378, App
305	27	56.2	485	9	US-10-508-263-22	Sequence 22, Appl	378	26	54.2	204	11	US-11-188-298-10553	Sequence 10553, A
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307	27	56.2	492	11	US-11-079-463-10294	Sequence 10294, A	380	26	54.2	213	11	US-11-096-568A-10205	Sequence 10205, A
308	27	56.2	494	11	US-11-094-519A-35	Sequence 35, Appl	381	26	54.2	217	11	US-11-087-099-4366	Sequence 4366, Ap
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310	27	56.2	513	11	US-11-098-686-10835	Sequence 10835, A	383	26	54.2	225	9	US-10-131-826A-282	Sequence 282, App
311	27	56.2	513	11	US-11-096-568A-29640	Sequence 29640, A	384	26	54.2	225	9	US-10-973-115B-282	Sequence 282, App
312	27	56.2	516	9	US-10-661-966-18	Sequence 18, Appl	385	26	54.2	225	9	US-10-137-873A-282	Sequence 282, App
313	27	56.2	561	8	US-10-505-928-163	Sequence 163, App	386	26	54.2	225	9	US-10-137-873A-282	Sequence 282, App

387	26	54.2	225	9	US-10-152-370-282	Sequence 282, App	460	26	54.2	365	11	US-11-188-298-2929	Sequence 2929, App
388	26	54.2	225	11	US-11-290-153-282	Sequence 283, App	461	26	54.2	365	11	US-11-188-298-7577	Sequence 7577, App
389	26	54.2	229	9	US-10-506-454-1521	Sequence 1521, App	462	26	54.2	366	11	US-11-152-669-161	Sequence 14, Appl
390	26	54.2	229	11	US-11-096-568A-13524	Sequence 13524, A	463	26	54.2	366	11	US-11-188-298-3602	Sequence 3602, App
391	26	54.2	232	11	US-11-096-568A-13523	Sequence 13523, A	464	26	54.2	368	11	US-11-188-298-2641	Sequence 2641, App
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418	26	54.2	325	9	US-10-506-454-1005	Sequence 1005, App	491	26	54.2	411	11	US-11-188-298-18719	Sequence 18719, A
419	26	54.2	325	11	US-11-096-568A-3823	Sequence 3823, App	492	26	54.2	416	11	US-11-188-298-13738	Sequence 13738, A
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421	26	54.2	327	11	US-11-096-568A-22705	Sequence 22705, A	494	26	54.2	417	11	US-11-109-156-2	Sequence 2, Appl1
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433	26	54.2	340	11	US-11-045-004-2010	Sequence 2010, App	506	26	54.2	435	11	US-11-188-298-18399	Sequence 18399, A
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439	26	54.2	351	11	US-11-096-568A-3326	Sequence 3326, App	512	26	54.2	455	11	US-11-188-298-441	Sequence 441, App
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540	26	502	9	US-10-219-061-4	Sequence 4, Appl	613	26	54.2	1464	11	US-11-124-678A-262	Sequence 262, Appl
541	26	502	9	US-10-219-062-4	Sequence 4, Appl	614	26	54.2	1493	11	US-11-183-136-26	Sequence 26, Appl
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549	26	502	11	US-11-230-995-3	Sequence 3718, Ap	622	26	54.2	2098	9	US-10-055-877-253	Sequence 253, App
550	26	502	11	US-11-072-512-3718	Sequence 14, Appl	623	26	54.2	2109	9	US-10-055-877-251	Sequence 251, App
551	26	502	9	US-10-912-971-14	Sequence 318, App	624	26	54.2	2204	9	US-10-495-083-8	Sequence 8, Appl
552	26	502	9	US-10-501-035-318	Sequence 318, App	625	26	54.2	2362	11	US-11-096-568A-31778	Sequence 31778, A
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561	26	502	11	US-10-087-099-6955	Sequence 6955, Ap	634	26	54.2	16	9	US-10-895-064-684	Sequence 684, App
562	26	502	9	US-10-455-772-456	Sequence 456, App	635	26	54.2	16	9	US-10-129-711-684	Sequence 711, App
563	26	502	9	US-10-455-772-452	Sequence 452, App	636	26	54.2	16	11	US-11-129-711-685	Sequence 182, App
564	26	502	9	US-10-506-454-152	Sequence 152, App	637	26	54.2	20	9	US-10-939-890-182	Sequence 91, Appl
565	26	502	11	US-11-072-512-3779	Sequence 3779, Ap	638	26	54.2	20	11	US-11-069-642-91	Sequence 12, Appl
566	26	502	11	US-11-087-099-7090	Sequence 7090, Ap	639	26	54.2	24	11	US-11-116-682-12	Sequence 330, App
567	26	502	11	US-11-087-099-4680	Sequence 4680, Ap	640	26	54.2	26	9	US-10-939-890-330	Sequence 326, App
568	26	502	11	US-11-043-788-376	Sequence 376, App	641	26	54.2	26	9	US-10-137-873A-84	Sequence 84, Appl
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571	26	502	11	US-11-096-568A-14616	Sequence 14616, A	644	26	54.2	26	9	US-10-939-890-336	Sequence 3655, Ap
572	26	502	11	US-11-043-788-374	Sequence 374, App	645	26	54.2	26	9	US-11-096-568A-27306	Sequence 27306, A
573	26	502	11	US-11-210-471-4	Sequence 4, Appl	646	26	54.2	42	11	US-11-087-099-7535	Sequence 7535, Ap
574	26	502	11	US-11-210-471-4	Sequence 10, Appl	647	26	54.2	53	11	US-11-087-099-9068	Sequence 63, Appl
575	26	502	11	US-11-043-788-375	Sequence 375, App	648	26	54.2	62	9	US-10-506-454-63	Sequence 9953, Ap
576	26	502	11	US-11-050-857-368	Sequence 388, App	649	26	54.2	79	11	US-11-096-568A-9953	Sequence 84, Appl
577	26	502	9	US-10-724-598-34	Sequence 34, Appl	650	26	54.2	80	9	US-10-137-873A-84	Sequence 84, Appl
578	26	502	11	US-11-050-857-384	Sequence 384, App	651	26	54.2	80	9	US-10-152-370-84	Sequence 3679, Ap
579	26	502	10	US-11-301-924-4	Sequence 4, Appl	652	26	54.2	80	9	US-11-096-568A-3679	Sequence 84, Appl
580	26	502	11	US-11-127-877-72	Sequence 72, Appl	653	26	54.2	80	11	US-11-290-153-84	Sequence 84, Appl
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582	26	502	11	US-11-188-298-3720	Sequence 3720, Ap	655	26	54.2	83	11	US-11-087-099-5230	Sequence 934, App
583	26	502	9	US-10-455-772-446	Sequence 446, App	656	26	54.2	93	11	US-11-000-463-934	Sequence 21911, A
584	26	502	6	US-11-188-298-14308	Sequence 14308, A	657	26	54.2	99	11	US-11-096-568A-21911	Sequence 5377, App
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586	26	502	9	US-10-455-772-450	Sequence 450, App	659	26	54.2	108	11	US-11-096-568A-13268	Sequence 656, App
587	26	502	11	US-11-052-554A-217	Sequence 217, App	660	26	54.2	109	11	US-09-978-360A-656	Sequence 9058, Ap
588	26	502	7	US-10-505-928-644	Sequence 644, App	661	26	54.2	113	7	US-11-087-099-9058	Sequence 462, App
589	26	502	8	US-10-511-937-2606	Sequence 2606, App	662	26	54.2	113	11	US-11-000-463-962	Sequence 21061, A
590	26	502	11	US-11-043-788-367	Sequence 368, App	663	26	54.2	113	11	US-11-096-568A-21061	Sequence 805, App
591	26	502	11	US-11-043-788-368	Sequence 373, App	664	26	54.2	124	11	US-11-172-740-805	Sequence 8920, Ap
592	26	502	11	US-11-043-788-373	Sequence 372, App	665	26	54.2	127	11	US-11-172-740-807	Sequence 8920, Ap
593	26	502	11	US-11-043-788-372	Sequence 18, Appl	666	26	54.2	128	11	US-11-096-568A-8920	Sequence 7104, Ap
594	26	502	11	US-11-197-133A-18	Sequence 7119, Ap	667	26	54.2	131	11	US-10-986-405-376	Sequence 376, App
595	26	502	11	US-11-079-463-7119	Sequence 370, App	668	26	54.2	132	9	US-10-986-405-376	Sequence 35, Appl
596	26	502	11	US-11-043-788-370	Sequence 5227, Ap	669	26	54.2	132	11	US-11-098-666-335	Sequence 10330, A
597	26	502	11	US-11-079-463-5227	Sequence 358, App	670	26	54.2	133	11	US-11-098-666-335	Sequence 744, App
598	26	502	9	US-10-131-826A-358	Sequence 358, App	671	26	54.2	136	7	US-09-978-360A-744	Sequence 8919, Ap
599	26	502	9	US-10-973-115B-358	Sequence 496, App	672	26	54.2	139	11	US-11-096-568A-8919	Sequence 8921, Ap
600	26	502	9	US-10-216-161A-496	Sequence 358, App	673	26	54.2	141	9	US-10-484-516-20	Sequence 2528, Ap
601	26	502	9	US-10-137-873A-358	Sequence 358, App	674	26	54.2	142	8	US-10-511-937-2528	Sequence 6504, Ap
602	26	502	11	US-10-152-370-358	Sequence 358, App	675	26	54.2	142	11	US-11-087-099-6504	Sequence 306, App
603	26	502	11	US-11-290-153-358	Sequence 144, App	676	26	54.2	142	11		
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682	25	52.1	155	9	US-10-793-626-1642	Sequence 1642, Ap	755	25	52.1	239	11	US-11-218-880-14	Sequence 14, Appl
683	25	52.1	155	11	US-11-207-078-227	Sequence 227, Ap	756	25	52.1	239	11	US-11-218-880-15	Sequence 15, Appl
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686	25	52.1	157	11	US-11-018-868-186	Sequence 186, Ap	759	25	52.1	241	11	US-11-175-590-289	Sequence 289, Ap
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881	25	52.1	378	11	US-11-087-099-11797	Sequence 11797, A	954	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
882	25	52.1	379	11	US-11-096-568A-14738	Sequence 14738, A	955	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
883	25	52.1	380	9	US-10-525-674-28	Sequence 28, Appl	956	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
884	25	52.1	384	11	US-11-096-568A-23002	Sequence 23002, A	957	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
885	25	52.1	385	11	US-11-087-099-7764	Sequence 7764, Ap	958	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
886	25	52.1	386	11	US-11-096-568A-11904	Sequence 11904, A	959	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
887	25	52.1	387	11	US-11-096-568A-19075	Sequence 19075, A	960	25	52.1	438	11	US-11-069-642-92	Sequence 92, Appl
888	25	52.1	389	11	US-11-088-686-1	Sequence 1, Appl1	961	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
889	25	52.1	389	11	US-11-088-686-5	Sequence 5, Appl1	962	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
890	25	52.1	389	11	US-11-088-686-7	Sequence 7, Appl1	963	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
891	25	52.1	389	11	US-11-088-686-9	Sequence 9, Appl1	964	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
892	25	52.1	389	11	US-11-088-686-11	Sequence 11, Appl	965	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
893	25	52.1	389	11	US-11-088-686-13	Sequence 13, Appl	966	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
894	25	52.1	390	9	US-10-485-517-235	Sequence 235, App	967	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
895	25	52.1	391	11	US-11-188-298-13048	Sequence 13048, A	968	25	52.1	442	9	US-10-506-454-1316	Sequence 4499, Ap
896	25	52.1	394	11	US-11-087-099-2746	Sequence 2746, Ap	969	25	52.1	462	11	US-11-188-298-10987	Sequence 10987, A
897	25	52.1	396	11	US-11-087-099-4329	Sequence 4329, Ap	970	25	52.1	462	11	US-11-188-298-16342	Sequence 16342, A

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971 25 52.1 465 11 US-11-087-039-6870 Sequence 6870, Ap
972 25 52.1 465 11 US-11-087-039-6865 Sequence 8665, Ap
973 25 52.1 465 11 US-11-188-238-19056 Sequence 19056, A
974 25 52.1 466 11 US-11-188-238-14612 Sequence 14612, A
975 25 52.1 469 9 US-10-793-626-2204 Sequence 2204, Ap
976 25 52.1 469 11 US-11-188-238-21184 Sequence 21184, A
977 25 52.1 473 11 US-11-074-176-338 Sequence 338, App
978 25 52.1 473 11 US-11-188-238-1021 Sequence 1021, Ap
979 25 52.1 476 11 US-11-024-959-232 Sequence 232, App
980 25 52.1 476 11 US-11-264-728-2 Sequence 2, Appli
981 25 52.1 478 9 US-10-330-773-861 Sequence 861, App
982 25 52.1 478 11 US-11-087-039-8625 Sequence 8625, A
983 25 52.1 478 11 US-11-188-238-19025 Sequence 19025, A
984 25 52.1 483 11 US-11-208-303-1 Sequence 1, Appli
985 25 52.1 484 11 US-11-126-313-24 Sequence 24, Appli
986 25 52.1 485 9 US-10-506-454-1364 Sequence 1364, Ap
987 25 52.1 485 11 US-11-087-039-398 Sequence 398, App
988 25 52.1 487 11 US-11-074-176-130 Sequence 130, App
989 25 52.1 491 11 US-11-188-238-18885 Sequence 18885, A
990 25 52.1 492 11 US-11-072-512-1974 Sequence 1974, Ap
991 25 52.1 492 11 US-11-072-512-3223 Sequence 3223, Ap
992 25 52.1 493 11 US-11-087-039-2109 Sequence 2109, Ap
993 25 52.1 493 11 US-11-087-039-6629 Sequence 6629, Ap
994 25 52.1 494 11 US-11-087-039-12046 Sequence 12046, A
995 25 52.1 495 11 US-11-066-568A-7940 Sequence 7940, Ap
996 25 52.1 496 11 US-11-188-238-16362 Sequence 16362, A
997 25 52.1 497 11 US-11-188-238-14336 Sequence 14336, A
998 25 52.1 498 11 US-11-188-238-14336 Sequence 2937, App
999 25 52.1 499 11 US-11-024-959-293 Sequence 293, App
1000 25 52.1 500 11 US-11-120-308-138 Sequence 138, App
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ALIGNMENTS

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RESULT 1
US-10-530-061-481
; Sequence 481, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: US/10/530,061
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: Patentin version 3.3
; SEQ ID NO 481
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-481

Query Match 100.0%; Score 48; DB 9; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.00091;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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QY 1 AMFODPOER 9
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Db 3 AMFODPOER 11

RESULT 2
US-10-530-061-482
; Sequence 482, Application US/10530061
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; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: US/10/530,061
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: Patentin version 3.3
; SEQ ID NO 482
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-482

Query Match 100.0%; Score 48; DB 9; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.00091;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 3
US-10-530-061-1702
; Sequence 1702, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: Patentin version 3.3
; SEQ ID NO 1702
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-1702

Query Match 100.0%; Score 48; DB 9; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.0013;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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QY 1 AMFODPOER 9
    |||||
Db 3 AMFODPOER 11

RESULT 4
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1
; GENERAL INFORMATION:
; APPLICANT: HealthBanks Biotech CO. LTD.
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/ TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
/ FILE REFERENCE: P7819/0613
/ CURRENT APPLICATION NUMBER: US/11/206,138
/ CURRENT FILING DATE: 2005-08-18
/ NUMBER OF SEQ ID NOS: 14
/ SOFTWARE: PatentIn version 3.3
/ SEQ ID NO 3
/ LENGTH: 158
/ TYPE: PRT
/ ORGANISM: Human papillomavirus type 16
US-11-206-138-3

Query Match 100.0%; Score 48; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.015;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 7 AMFODPOER 15

RESULT 5
US-11-192-923A-2
/ Sequence 2, Application US/11192923A
/ Publication No. US20060018928A1
/ GENERAL INFORMATION:
/ APPLICANT: PANG, XIAOMU
/ TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
/ FILE REFERENCE: 116620-003
/ CURRENT APPLICATION NUMBER: US/11/192,923A
/ CURRENT FILING DATE: 2005-07-29
/ PRIOR APPLICATION NUMBER: CN 03115272.4
/ PRIOR FILING DATE: 2003-01-30
/ PRIOR APPLICATION NUMBER: CN 03115273.2
/ PRIOR FILING DATE: 2003-01-30
/ NUMBER OF SEQ ID NOS: 45
/ SOFTWARE: PatentIn Ver. 3.3
/ SEQ ID NO 2
/ LENGTH: 256
/ TYPE: PRT
/ ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 48; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 105 AMFODPOER 113

RESULT 6
US-10-530-061-483
/ Sequence 483, Application US/10530061
/ Publication No. US2006007945A1
/ GENERAL INFORMATION:
/ APPLICANT: SIDNEY, JOHN
/ APPLICANT: SOUTHWOOD, SCOTT
/ APPLICANT: SETTE, ALESSANDRO
/ TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
/ FILE REFERENCE: 2060.033US02/EKS/M-M
/ CURRENT APPLICATION NUMBER: US/10/530,061
/ CURRENT FILING DATE: 2005-04-04
/ PRIOR APPLICATION NUMBER: PCT/US03/31308
/ PRIOR FILING DATE: 2003-10-03
/ PRIOR APPLICATION NUMBER: 60/416,207
/ PRIOR FILING DATE: 2002-10-03
/ PRIOR APPLICATION NUMBER: 60/417,269
/ PRIOR FILING DATE: 2002-10-08
/ NUMBER OF SEQ ID NOS: 2503
/ SOFTWARE: PatentIn version 3.3

/ SEQ ID NO 483
/ LENGTH: 11
/ TYPE: PRT
/ ORGANISM: Human papillomavirus
US-10-530-061-483

Query Match 91.7%; Score 44; DB 9; Length 11;
Best Local Similarity 100.0%; Pred. No. 0.0059;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 1 MFODPOER 8

RESULT 7
US-10-530-253-13
/ Sequence 13, Application US/10530253
/ Publication No. US20060014926A1
/ GENERAL INFORMATION:
/ APPLICANT: Casabetti, Maria C.
/ APPLICANT: Smith, Larry
/ APPLICANT: Jeffrey K. Pullen
/ APPLICANT: Susan P. McElhinney
/ TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
/ FILE REFERENCE: 00630/100M137-US2
/ CURRENT APPLICATION NUMBER: US/10/530,253
/ CURRENT FILING DATE: 2005-04-04
/ PRIOR APPLICATION NUMBER: PCT/US2003/031726
/ PRIOR FILING DATE: 2003-10-02
/ PRIOR APPLICATION NUMBER: US 60/415,929
/ PRIOR FILING DATE: 2002-10-03
/ NUMBER OF SEQ ID NOS: 65
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 13
/ LENGTH: 151
/ TYPE: PRT
/ ORGANISM: Human papillomavirus type 16
US-10-530-253-13

Query Match 91.7%; Score 44; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.092;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 1 MFODPOER 8

RESULT 8
US-10-530-253-1
/ Sequence 1, Application US/10530253
/ Publication No. US20060014926A1
/ GENERAL INFORMATION:
/ APPLICANT: Casabetti, Maria C.
/ APPLICANT: Smith, Larry
/ APPLICANT: Jeffrey K. Pullen
/ APPLICANT: Susan P. McElhinney
/ TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
/ FILE REFERENCE: 00630/100M137-US2
/ CURRENT APPLICATION NUMBER: US/10/530,253
/ CURRENT FILING DATE: 2005-04-04
/ PRIOR APPLICATION NUMBER: PCT/US2003/031726
/ PRIOR FILING DATE: 2003-10-02
/ PRIOR APPLICATION NUMBER: US 60/415,929
/ PRIOR FILING DATE: 2002-10-03
/ NUMBER OF SEQ ID NOS: 65
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 1
/ LENGTH: 248
/ TYPE: PRT
/ ORGANISM: Human papillomavirus type 16
US-10-530-253-1

Query Match 91.7%; Score 44; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 1 MFODPOER 8

RESULT 9
US-10-530-253-3

; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 91.7%; Score 44; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 1 MFODPOER 8

RESULT 10
US-10-530-253-5

; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 91.7%; Score 44; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
Db 1 MFODPOER 8

RESULT 11
US-10-530-253-7

; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 81.2%; Score 39; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
Db 99 FODPOER 105

RESULT 12
US-10-530-253-9

; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 81.2%; Score 39; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPOER 9
Db 99 FODPOER 105

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RESULT 13
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhiney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match      81.2%; Score 39; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 FQDPOER 9
Db 99 FQDPOER 105

RESULT 14
US-10-530-253-18
; Sequence 18, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhiney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 35
US-10-530-253-18

Query Match      79.2%; Score 38; DB 9; Length 149;
Best Local Similarity 87.5%; Pred. No. 1.5;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 1 MFQDPOER 8

RESULT 15
US-11-079-463-7932
; Sequence 7932, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
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; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRP
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/126,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 7932
; LENGTH: 567
; TYPE: PRT
; ORGANISM: B. fragilis
US-11-079-463-7932

Query Match      75.0%; Score 36; DB 11; Length 567;
Best Local Similarity 55.6%; Pred. No. 15;
Matches 5; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFQDPOER 9
Db 298 SMYQDPOER 306

RESULT 16
US-11-096-568A-18478
; Sequence 18478, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; PRIOR FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 18478
; LENGTH: 144
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(144)
; OTHER INFORMATION: Ceres Seq. ID no. 12365374
US-11-096-568A-18478

Query Match      66.7%; Score 32; DB 11; Length 144;
Best Local Similarity 62.5%; Pred. No. 24;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPOER 9
Db 122 LMQDPOER 129

RESULT 17
US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhiney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
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PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Patencin version 3.1
SEQ ID NO 26
LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 68
US-10-530-253-26

Query Match 66.7%; Score 32; DB 9; Length 158;
Best Local Similarity 55.6%; Pred. No. 26;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 2 ALFHNPEER 10

RESULT 18
US-11-096-568A-18476
Sequence 18476, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 18476
LENGTH: 167
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc.feature
LOCATION: (1)-(167)
OTHER INFORMATION: Ceres Seq. ID no. 12365372
US-11-096-568A-18476

Query Match 66.7%; Score 32; DB 11; Length 167;
Best Local Similarity 62.5%; Pred. No. 28;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 145 LWODPOEQ 152

RESULT 19
US-11-188-298-16639
Sequence 16639, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 16639
LENGTH: 288
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
NAME/KEY: unsure
LOCATION: (1)-(288)
OTHER INFORMATION: unsure at all Xaa locations
US-11-188-298-16639

Query Match 66.7%; Score 32; DB 11; Length 288;

Best Local Similarity 75.0%; Pred. No. 49;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 137 MFLNPOER 144

RESULT 20
US-10-505-928-327
Sequence 327, Application US/10505928
Publication No. US20060088532A1
GENERAL INFORMATION:
APPLICANT: Ludwig Institute for Cancer Research et al.
TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES
FILE REFERENCE: 28967/39178
CURRENT APPLICATION NUMBER: US/10/505,928
CURRENT FILING DATE: 2004-08-27
PRIOR APPLICATION NUMBER: US 60/363,019
PRIOR FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 866
SOFTWARE: Patencin 3.2
SEQ ID NO 327
LENGTH: 325
TYPE: PRT
ORGANISM: Homo sapiens
US-10-505-928-327

Query Match 66.7%; Score 32; DB 8; Length 325;
Best Local Similarity 62.5%; Pred. No. 56;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 157 VFEDPOER 164

RESULT 21
US-11-301-554-1816
Sequence 1816, Application US/11301554
Publication No. US20060088527A1
GENERAL INFORMATION:
APPLICANT: Henderson, Robert A.
APPLICANT: Wang, Tonglong
APPLICANT: Matanabe, Yoshihiro
APPLICANT: Kalos, Michael D.
APPLICANT: Sleach, Paul R.
APPLICANT: Johnson, Jeffrey C.
APPLICANT: Retter, Marc W.
APPLICANT: Durham, Margarita
APPLICANT: Carter, Darriek
APPLICANT: Fanger, Gary R.
APPLICANT: Vedvick, Thomas S.
APPLICANT: Bangur, Chaitanya S.
APPLICANT: McNabb, Andrea
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF LUNG CANCER
FILE REFERENCE: 210121.478C21
CURRENT APPLICATION NUMBER: US/11/301,554
CURRENT FILING DATE: 2005-12-13
PRIOR APPLICATION NUMBER: US 10/283,017
PRIOR FILING DATE: 2002-10-28
PRIOR APPLICATION NUMBER: US 10/113,872
PRIOR FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: US 10/017,754
PRIOR FILING DATE: 2001-10-29
PRIOR APPLICATION NUMBER: US 09/902,941
PRIOR FILING DATE: 2001-07-10
PRIOR APPLICATION NUMBER: US 09/849,626
PRIOR FILING DATE: 2001-05-03
PRIOR APPLICATION NUMBER: US 09/736,457
PRIOR FILING DATE: 2000-12-13
PRIOR APPLICATION NUMBER: US 09/702,705


```

; PRIOR FILING DATE: 2000-10-30
; PRIOR APPLICATION NUMBER: US 09/677,419
; PRIOR FILING DATE: 2000-10-06
; PRIOR APPLICATION NUMBER: US 09/671,325
; PRIOR FILING DATE: 2000-09-26
; PRIOR APPLICATION NUMBER: US 09/658,824
; PRIOR FILING DATE: 2000-09-08
; Remaining Prior Application data removed - See file wrapper or PALM.
; NUMBER OF SEQ ID NOS: 2157
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO: 1816
; LENGTH: 325
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-301-554-1816

```

```

Query Match      66.7%; Score 32; DB 10; Length 325;
Best Local Similarity 62.5%; Pred. No. 56;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      2 MFODPOER 9
DB      157 VFEDPORR 164

```

```

RESULT 22
US-11-188-298-13197
; Sequence 13197, Application US/11188298
; Publication No. US2006005522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO: 13197
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Zea mays
US-11-188-298-13197

```

```

Query Match      66.7%; Score 32; DB 11; Length 329;
Best Local Similarity 62.5%; Pred. No. 56;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 AMFODPOE 8
DB      315 AMFEDPSQ 322

```

```

RESULT 23
US-10-821-234-1163
; Sequence 1163, Application US/10821234
; Publication No. US20050255114A1
; GENERAL INFORMATION:
; APPLICANT: Labat, Ivan
; APPLICANT: Stache-Crain, Birgit
; APPLICANT: Andarmant, Susan
; APPLICANT: Tang, Y. Tom
; TITLE OF INVENTION: Methods for Diagnosis and Treatment of Preeclampsia
; FILE REFERENCE: 821A
; CURRENT APPLICATION NUMBER: US/10/821,234
; CURRENT FILING DATE: 2004-04-07
; PRIOR APPLICATION NUMBER: US 60/462,047
; PRIOR FILING DATE: 2003-04-07
; NUMBER OF SEQ ID NOS: 1704
; SOFTWARE: pt SEQ_genes Version 1.0
; SEQ ID NO: 1163
; LENGTH: 399
; TYPE: PRT

```

```

; ORGANISM: Homo sapiens
US-10-821-234-1163

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```

Query Match      66.7%; Score 32; DB 9; Length 399;
Best Local Similarity 62.5%; Pred. No. 69;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      2 MFODPOER 9
DB      231 VFEDPORR 238

```

```

RESULT 24
US-11-096-568A-32147
; Sequence 32147, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO: 32147
; LENGTH: 1341
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(1341)
; OTHER INFORMATION: Ceres Seq. ID no. 13592292
US-11-096-568A-32147

```

```

Query Match      66.7%; Score 32; DB 11; Length 1341;
Best Local Similarity 62.5%; Pred. No. 2,5e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 AMFODPOE 8
DB      1085 AFEDPEE 1092

```

```

RESULT 25
US-11-096-568A-32146
; Sequence 32146, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO: 32146
; LENGTH: 1372
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(1372)
; OTHER INFORMATION: Ceres Seq. ID no. 13592291
US-11-096-568A-32146

```

```

Query Match      66.7%; Score 32; DB 11; Length 1372;
Best Local Similarity 62.5%; Pred. No. 2,5e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 AMFODPOE 8
DB      1116 AFEDPEE 1123

```

```
RESULT 26
US-11-096-568A-32145
; Sequence 32145, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 32145
; LENGTH: 1393
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(1393)
; OTHER INFORMATION: Ceres Seq. ID no. 13592290
US-11-096-568A-32145

Query Match          66.7%; Score 32; DB 11; Length 1393;
Best Local Similarity 62.5%; Pred. No. 2,6e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 AMFODPOE 8
|:|:|:|
DB      1137 AFEDPBE 1144

RESULT 27
US-10-530-253-39
; Sequence 39, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 39
; LENGTH: 152
; TYPE: PRT
; ORGANISM: Human papillomavirus
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (1)..(152)
; OTHER INFORMATION: where Xaa is any amino acid
US-10-530-253-39

Query Match          64.6%; Score 31; DB 9; Length 152;
Best Local Similarity 71.4%; Pred. No. 40;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      3 FODPOER 9
|:|:|:|
DB      4 FZDPXER 10

RESULT 28
US-11-096-568A-18941
; Sequence 18941, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 18941
; LENGTH: 445
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(445)
; OTHER INFORMATION: Ceres Seq. ID no. 12367925
US-11-096-568A-18941

Query Match          64.6%; Score 31; DB 11; Length 445;
Best Local Similarity 55.6%; Pred. No. 1,2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 AMFODPOER 9
|:|:|:|
DB      144 AVFEPXER 152

RESULT 29
US-10-491-468-5
; Sequence 5, Application US/10491468
; Publication No. US20060051836A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE CORPORATION; TANG, Y. Tom;
; APPLICANT: FORSYTHE, Ian J.; EMERLING, Brooke M.;
; APPLICANT: HAFALIA, April J.A.; YDE, Henry;
; APPLICANT: XU, Yuming; GIETZEN, Kimberly J.;
; APPLICANT: CHAWLA, Nandinder K.; BAUGHN, Mariah R.;
; APPLICANT: MARQUIS, Joseph P.; BECHA, Shanya D.;
; APPLICANT: KABLE, Amy E.; LALU, Preeti G.;
; APPLICANT: RICHARDSON, Thomas W.; LEE, Soo Y.;
; APPLICANT: LEE, Ernestine A.; TRAN, Bao;
; APPLICANT: WARREN, Bridget A.; LU, Dyoung Aina M.;
; APPLICANT: GURURAJAN, Rajagopal; SPRAGUE, William W.;
; APPLICANT: BLAKE, Julie J.; THANGAVELU, Kavitha;
; APPLICANT: SWARNAKAR, Anita; GORVAD, Ann E.;
; APPLICANT: GRIFFIN, Jennifer A.; LINDQUIST, Erika A.;
; APPLICANT: ELLIOTT, Vicki S.; ISON, Craig H.;
; APPLICANT: RAMKUMAR, Jayalaxmi
; TITLE OF INVENTION: MOLECULES FOR DISEASE DETECTION AND TREATMENT
; FILE REFERENCE: PF-1232 USN
; CURRENT APPLICATION NUMBER: US/10/491,468
; CURRENT FILING DATE: 2004-03-31
; PRIOR APPLICATION NUMBER: PCT/US02/32852
; PRIOR FILING DATE: 2002-10-10
; PRIOR APPLICATION NUMBER: US 60/328,944
; PRIOR FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: US 60/345,384
; PRIOR FILING DATE: 2001-10-26
; PRIOR APPLICATION NUMBER: US 60/343,880
; PRIOR FILING DATE: 2001-11-02
; PRIOR APPLICATION NUMBER: US 60/345,143
; PRIOR FILING DATE: 2001-11-09
; PRIOR APPLICATION NUMBER: US 60/332,430
; PRIOR FILING DATE: 2001-11-16
; NUMBER OF SEQ ID NOS: 96
; SOFTWARE: PERL Program
; SEQ ID NO 5
; LENGTH: 447
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
```

OTHER INFORMATION: Incyte ID No: 7644881CD1
US-10-491-468-5

Query Match 64.6%; Score 31; DB 9; Length 447;
Best Local Similarity 55.6%; Pred. No. 1.2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 36 ALFOPPREX 44

RESULT 30
US-11-096-568A-18940
; Sequence 18940, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 18940
; LENGTH: 470
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(470)
; OTHER INFORMATION: Ceres Seq. ID no. 12367924
US-11-096-568A-18940

Query Match 64.6%; Score 31; DB 11; Length 470;
Best Local Similarity 55.6%; Pred. No. 1.3e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 169 AVFEEPAER 177

RESULT 31
US-11-096-568A-18939
; Sequence 18939, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 18939
; LENGTH: 486
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(486)
; OTHER INFORMATION: Ceres Seq. ID no. 12367923
US-11-096-568A-18939

Query Match 64.6%; Score 31; DB 11; Length 486;
Best Local Similarity 55.6%; Pred. No. 1.4e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPOER 9
DB 185 AVFEEPAER 193

RESULT 32
US-10-530-253-17

; Sequence 17, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 17
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 33
US-10-530-253-17

Query Match 62.5%; Score 30; DB 9; Length 149;
Best Local Similarity 62.5%; Pred. No. 63;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 1 MFODTEEK 8

RESULT 33
US-10-530-253-24
; Sequence 24, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 24
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 58
US-10-530-253-24

Query Match 62.5%; Score 30; DB 9; Length 149;
Best Local Similarity 62.5%; Pred. No. 63;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFODPOER 9
DB 1 MFODABEK 8

RESULT 34
US-10-530-253-20
; Sequence 20, Application US/10530253
; Publication No. US20060014926A1

```

; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2003-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 45
; US-10-530-253-20

Query Match          62.5%; Score 30; DB 9; Length 158;
Best Local Similarity 55.6%; Pred. No. 66;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPQR 9
|:|:|:|
Db 2 ARFDPPQR 10

RESULT 35
; US-10-530-253-25
; Sequence 25, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2003-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 25
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Human papillomavirus type 59
; US-10-530-253-25

Query Match          62.5%; Score 30; DB 9; Length 160;
Best Local Similarity 55.6%; Pred. No. 67;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPQR 9
|:|:|:|
Db 2 ARFDPPQR 10

RESULT 36
; US-10-499-246-17
; Sequence 17, Application US/10499246
; Publication No. US20060053512A1
; GENERAL INFORMATION:
; APPLICANT: Bao, Xiaoming
; APPLICANT: Ohlrogge, John B.
; APPLICANT: Pollard, Michael R.
; TITLE OF INVENTION: Plant Cyclopropane Fatty Acid Synthase Genes, Proteins, and Uses
```

```

; TITLE OF INVENTION: Thereof
; FILE REFERENCE: DOW-MSU-07661
; CURRENT APPLICATION NUMBER: US/10/499,246
; CURRENT FILING DATE: 2004-06-17
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 17
; LENGTH: 293
; TYPE: PRT
; ORGANISM: Cyclopropane synthase
; US-10-499-246-17

Query Match          62.5%; Score 30; DB 9; Length 293;
Best Local Similarity 83.3%; Pred. No. 1,3e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODP 6
|:|:|:|
Db 30 ALFODP 35

RESULT 37
; US-10-499-246-18
; Sequence 18, Application US/10499246
; Publication No. US20060053512A1
; GENERAL INFORMATION:
; APPLICANT: Bao, Xiaoming
; APPLICANT: Ohlrogge, John B.
; APPLICANT: Pollard, Michael R.
; TITLE OF INVENTION: Plant Cyclopropane Fatty Acid Synthase Genes, Proteins, and Uses
; FILE REFERENCE: DOW-MSU-07661
; CURRENT APPLICATION NUMBER: US/10/499,246
; CURRENT FILING DATE: 2004-06-17
; NUMBER OF SEQ ID NOS: 35
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 18
; LENGTH: 301
; TYPE: PRT
; ORGANISM: Cyclopropane synthase
; US-10-499-246-18

Query Match          62.5%; Score 30; DB 9; Length 301;
Best Local Similarity 83.3%; Pred. No. 1,3e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODP 6
|:|:|:|
Db 33 ALFODP 38

RESULT 38
; US-10-506-454-1478
; Sequence 1478, Application US/10506454
; Publication No. US20060068386A1
; GENERAL INFORMATION:
; APPLICANT: Slesarev, Alexi I
; APPLICANT: Meznevaya, Katja V
; APPLICANT: Polushin, Nikolai N
; APPLICANT: Shcherbina, Olga V
; APPLICANT: Shakhova, Vera V
; APPLICANT: Malykh, Andrei G
; APPLICANT: Kozlyavkin, Sergei A
; TITLE OF INVENTION: The Complete Genome and Protein Sequences of the Hyperthermophile
; TITLE OF INVENTION: Methanopyrus Kandleri AV19 and Monophyly of Archaeal Methanogens
; FILE REFERENCE: FID001
; CURRENT APPLICATION NUMBER: US/10/506,454
; CURRENT FILING DATE: 2004-08-31
; PRIOR APPLICATION NUMBER: PCT/US03/06664
; PRIOR FILING DATE: 2003-03-04
; PRIOR APPLICATION NUMBER: 60/361,742
; PRIOR FILING DATE: 2002-03-04
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NUMBER OF SEQ ID NOS: 1722
SOFTWARE: PatentIn version 3.2
SEQ ID NO 1478
LENGTH: 309
TYPE: PRT
ORGANISM: Methanopyrus kandleri
US-10-506-454-1478

Query Match 62.5%; Score 30; DB 9; Length 309;
Best Local Similarity 71.4%; Pred. No. 1.3e+02;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 MFQDPQE 8
DB 234 MFDPDE 240

RESULT 39
US-10-793-626-3238
Sequence 3238, Application US/10793626
Publication No. US20050255478A1
GENERAL INFORMATION:
APPLICANT: KIMBERLY, WILLIAM JOHN
TITLE OF INVENTION: STAPHYLOCOCCUS EPIDERMIDIS NUCLEIC ACIDS AND PROTEINS
FILE REFERENCE: PU3480US
CURRENT APPLICATION NUMBER: US/10/793,626
CURRENT FILING DATE: 2004-03-04
PRIOR APPLICATION NUMBER: 60/164,258
PRIOR FILING DATE: 1999-11-09
NUMBER OF SEQ ID NOS: 4472
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3238
LENGTH: 326
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
TITLE OF INVENTION: Description of Artificial Sequence: synthetic
OTHER INFORMATION: amino acid sequence
US-10-793-626-3238

Query Match 62.5%; Score 30; DB 9; Length 326;
Best Local Similarity 75.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFQDPOE 8
DB 109 AMFQDPLE 116

RESULT 40
US-11-082-389-132
Sequence 132, Application US/11082389
Publication No. US2005024935A1
GENERAL INFORMATION:
APPLICANT: Pompejus, Markus
APPLICANT: Krogger, Burkhard
APPLICANT: Schröder, Hartwig
APPLICANT: Zelder, Oskar
APPLICANT: Habedanauer, Gregor
TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING PROTEINS
TITLE OF INVENTION: INVOLVED IN MEMBRANE SYNTHESIS AND MEMBRANE
FILE REFERENCE: BGI-131CPCN
CURRENT APPLICATION NUMBER: US/11/082,389
CURRENT FILING DATE: 2005-03-16
PRIOR APPLICATION NUMBER: US 09/603024
PRIOR FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: US 60/141031
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: US 60/143262
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: US 60/151281
PRIOR FILING DATE: 1999-08-27

PRIOR APPLICATION NUMBER: DE 19930487.4
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: DE 19930489.0
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: DE 19931549.3
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19931550.7
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19932134.5
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: DE 19941379.7
PRIOR FILING DATE: 1999-08-31
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 446
SEQ ID NO 132
LENGTH: 365
TYPE: PRT
ORGANISM: Corynebacterium glutamicum
US-11-082-389-132

Query Match 62.5%; Score 30; DB 11; Length 365;
Best Local Similarity 83.3%; Pred. No. 1.6e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFQDPQ 7
DB 202 VFQDPQ 207

RESULT 41
US-11-096-568A-29694
Sequence 29694, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 29694
LENGTH: 367
TYPE: PRT
ORGANISM: Arabidopsis thaliana
FEATURE:
NAME/KEY: misc.feature
LOCATION: (1)..(367)
OTHER INFORMATION: Ceres Seq. ID no. 4925266
US-11-096-568A-29694

Query Match 62.5%; Score 30; DB 11; Length 367;
Best Local Similarity 55.6%; Pred. No. 1.6e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFQDPOE 9
DB 235 AAFDDEEQ 243

RESULT 42
US-11-096-568A-29693
Sequence 29693, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 29693

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; LENGTH: 380
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(380)
; OTHER INFORMATION: Ceres Seq. ID no. 4925265
US-11-096-568A-29693

Query Match          62.5%; Score 30; DB 11; Length 380;
Best Local Similarity 55.6%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPQER 9
Db 248 AAFDDPEEQ 256

RESULT 43
US-11-096-568A-29692
; Sequence 29692, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 29692
; LENGTH: 416
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(416)
; OTHER INFORMATION: Ceres Seq. ID no. 4925264
US-11-096-568A-29692

Query Match          62.5%; Score 30; DB 11; Length 416;
Best Local Similarity 55.6%; Pred. No. 1.8e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 AMFODPQER 9
Db 284 AAFDDPEEQ 292

RESULT 44
US-11-082-389-130
; Sequence 130, Application US/11082389
; Publication No. US20050244935A1
; GENERAL INFORMATION:
; APPLICANT: Pompeius, Markus
; APPLICANT: Kroger, Burkhard
; APPLICANT: Schroder, Hartwig
; APPLICANT: Zelder, Oskar
; APPLICANT: Habehauer, Gregor
; TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING PROTEINS
; TITLE OF INVENTION: INVOLVED IN MEMBRANE SYNTHESIS AND MEMBRANE
; TITLE OF INVENTION: TRANSPORT
; FILE REFERENCE: BGI-131PCPN
; CURRENT APPLICATION NUMBER: US/11/082,389
; CURRENT FILING DATE: 2005-03-16
; PRIOR APPLICATION NUMBER: US 09/603024
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: US 60/141031
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: US 60/143262
; PRIOR FILING DATE: 1999-07-09
; PRIOR APPLICATION NUMBER: US 60/151281
; PRIOR FILING DATE: 1999-08-27
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; PRIOR APPLICATION NUMBER: DE 19930487.4
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: DE 19930489.0
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: DE 19931549.3
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19931550.7
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19932134.5
; PRIOR FILING DATE: 1999-07-09
; PRIOR APPLICATION NUMBER: DE 19941379.7
; PRIOR FILING DATE: 1999-08-31
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 446
; SEQ ID NO 130
; LENGTH: 479
; TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
US-11-082-389-130

Query Match          62.5%; Score 30; DB 11; Length 479;
Best Local Similarity 83.3%; Pred. No. 2.1e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 MFODPQ 7
Db 316 VFODPQ 321

RESULT 45
US-11-188-298-16827
; Sequence 16827, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16827
; LENGTH: 510
; TYPE: PRT
; ORGANISM: Oryza sativa (japonica cultivar-group)
US-11-188-298-16827

Query Match          62.5%; Score 30; DB 11; Length 510;
Best Local Similarity 83.3%; Pred. No. 2.3e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 FODPQ 8
Db 411 FODPQ 416

RESULT 46
US-11-079-463-6591
; Sequence 6591, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRA
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
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SEQ ID NO 6591
LENGTH: 533
TYPE: PRT
ORGANISM: B.fragilis
US-11-079-463-6591

Query Match 62.5%; Score 30; DB 11; Length 533;
Best Local Similarity 85.7%; Pred. No. 2.4e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 AMFODPQ 7
| | | | |
Db 64 AMFODKQ 70

RESULT 47
US-11-188-298-20601
Sequence 20601, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 20601
LENGTH: 543
TYPE: PRT
ORGANISM: Trichoderma harzianum
US-11-188-298-20601

Query Match 62.5%; Score 30; DB 11; Length 543;
Best Local Similarity 83.3%; Pred. No. 2.4e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODP 6
| | | | |
Db 488 AMFEDP 493

RESULT 48
US-11-188-298-22276
Sequence 22276, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 22276
LENGTH: 548
TYPE: PRT
ORGANISM: Hypocrea jecorina
US-11-188-298-22276

Query Match 62.5%; Score 30; DB 11; Length 548;
Best Local Similarity 83.3%; Pred. No. 2.4e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 AMFODP 6
| | | | |
Db 488 AMFEDP 493

RESULT 49
US-11-087-099-2496

Sequence 2496, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 2496
LENGTH: 601
TYPE: PRT
ORGANISM: Oryza sativa
FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(601)
OTHER INFORMATION: unsure at all Xaa locations
US-11-087-099-2496

Query Match 62.5%; Score 30; DB 11; Length 601;
Best Local Similarity 66.7%; Pred. No. 2.7e+02;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 AMFODPQER 9
| | | | |
Db 71 AMSQDPSSR 79

RESULT 50
US-11-087-099-11892
Sequence 11892, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 11892
LENGTH: 648
TYPE: PRT
ORGANISM: Oryza sativa
FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(648)
OTHER INFORMATION: unsure at all Xaa locations
US-11-087-099-11892

Query Match 62.5%; Score 30; DB 11; Length 648;
Best Local Similarity 66.7%; Pred. No. 2.9e+02;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 AMFODPQER 9
| | | | |
Db 71 AMSQDPSSR 79

Search completed: May 5, 2006, 08:56:11
Job time : 10 secs

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GenCore version 5.1.7
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OM protein - protein search, using bw model

Run on: May 5, 2006, 05:36:41 / Search time 20.9 Seconds
(without alignments)
35.602 Million cell updates/sec

Title: US-08-170-344-2
Perfect score: 48
Sequence: 1 KLPOLCTEL 9

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :
1: /cgn2_6/ptodata/1/1aa/5.COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/6.COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/H.COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/PCITUS.COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/RE.COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	48	100.0	9	1	US-08-787-547-101
2	48	100.0	9	2	US-09-601-723-274
3	48	100.0	20	1	US-08-934-915-159
4	48	100.0	30	1	US-08-363-586-4
5	48	100.0	30	2	US-09-980-523A-4
6	48	100.0	151	2	US-09-701-080C-18
7	48	100.0	158	2	US-09-980-523A-2
8	48	100.0	162	1	US-08-316-239B-3
9	48	100.0	162	1	US-08-316-239B-4
10	48	100.0	172	2	US-08-860-165-14
11	48	100.0	172	2	US-09-359-382-14
12	48	100.0	182	1	US-08-117-083-10
13	48	100.0	243	2	US-09-462-993-1
14	48	100.0	266	2	US-08-860-165-10
15	48	100.0	266	2	US-09-359-382-10
16	48	100.0	266	2	US-09-367-309A-1
17	48	100.0	273	2	US-09-485-885-4
18	48	100.0	292	2	US-09-485-885-10
19	48	100.0	371	2	US-09-485-885-6
20	48	100.0	390	2	US-09-485-885-14
21	43	89.6	32	1	US-08-466-285-2
22	43	89.6	32	2	US-08-164-768-2
23	43	89.6	158	2	US-08-247-904B-10
24	43	89.6	158	2	US-08-767-942A-19
25	43	89.6	271	1	US-08-117-083-14
26	43	89.6	278	1	US-09-485-885-21
27	43	89.6	383	2	US-09-485-885-23

28	72.9	211	2	US-09-543-681A-8111	Sequence 8111, App
29	70.8	15	1	US-07-909-122-2	Sequence 2, Appl
30	70.8	15	2	US-08-075-541D-52	Sequence 52, Appl
31	70.8	23	1	US-08-363-586-3	Sequence 3, Appl
32	70.8	161	2	US-09-673-395A-248	Sequence 248, App
33	70.8	341	2	US-08-481-968A-11	Sequence 11, Appl
34	70.8	341	2	US-08-154-712B-11	Sequence 11, Appl
35	70.8	341	2	US-09-947-925A-11	Sequence 11, Appl
36	70.8	504	2	US-09-198-452A-1036	Sequence 1036, App
37	70.8	504	2	US-09-438-185A-966	Sequence 966, App
38	70.8	522	2	US-09-949-016-10663	Sequence 10663, A
39	70.8	728	2	US-09-747-259-18	Sequence 18, Appl
40	70.8	728	2	US-09-816-744-18	Sequence 18, Appl
41	70.8	728	2	US-10-104-047-3399	Sequence 3399, App
42	68.8	163	2	US-09-252-991A-3145	Sequence 3145, A
43	68.8	304	2	US-09-248-796A-16504	Sequence 16504, A
44	68.8	308	2	US-09-369-247-60	Sequence 60, Appl
45	68.8	308	2	US-10-062-548-60	Sequence 60, Appl
46	68.8	384	2	US-10-104-047-3188	Sequence 3188, App
47	68.8	476	2	US-09-543-681A-7117	Sequence 7117, App
48	68.8	536	1	US-08-551-211-3	Sequence 3, Appl
49	68.8	620	2	US-09-673-198-1	Sequence 1, Appl
50	68.8	626	2	US-09-489-039A-13113	Sequence 13113, A
51	68.8	645	2	US-09-543-681A-6140	Sequence 6140, App
52	66.7	72	2	US-09-583-110-4207	Sequence 4207, App
53	66.7	73	2	US-09-248-796A-21511	Sequence 21511, A
54	66.7	86	2	US-09-348-796A-15475	Sequence 15475, A
55	66.7	91	2	US-09-342-647-20	Sequence 20, Appl
56	66.7	197	2	US-09-270-767-33420	Sequence 33420, A
57	66.7	203	2	US-09-270-767-48637	Sequence 48637, A
58	66.7	203	2	US-09-328-352-4681	Sequence 4681, App
59	66.7	236	2	US-09-634-137-32	Sequence 32, Appl
60	66.7	270	2	US-09-107-532A-4778	Sequence 4778, App
61	66.7	327	2	US-09-574-681A-4435	Sequence 4435, App
62	66.7	357	2	US-09-574-942-4	Sequence 4, Appl
63	66.7	357	2	US-09-949-434-4	Sequence 4, Appl
64	66.7	357	2	US-10-240-709-4	Sequence 4, Appl
65	66.7	361	2	US-08-807-784B-3	Sequence 3, Appl
66	66.7	505	2	US-09-270-767-43800	Sequence 43800, A
67	66.7	1323	2	US-09-489-039A-13945	Sequence 13945, A
68	66.7	1326	2	US-09-253-991A-17932	Sequence 17932, A
69	66.7	1385	2	US-09-827-998-16	Sequence 16, Appl
70	66.7	1690	2	US-09-595-664B-39	Sequence 39, Appl
71	66.7	1690	2	US-09-949-016-6474	Sequence 6474, App
72	66.7	1695	2	US-09-949-016-9887	Sequence 9887, App
73	66.7	1791	2	US-09-827-998-10	Sequence 10, Appl
74	66.7	1791	2	US-09-854-133-425	Sequence 425, App
75	66.7	4019	2	US-08-319-387-7	Sequence 7, Appl
76	64.6	70	2	US-09-430-029-2	Sequence 2, Appl
77	64.6	76	2	US-09-513-999C-4685	Sequence 4685, App
78	64.6	82	2	US-09-198-432A-1103	Sequence 1103, App
79	64.6	147	2	US-09-513-999C-4238	Sequence 4238, App
80	64.6	165	2	US-09-621-976-5263	Sequence 5263, App
81	64.6	170	1	US-08-650-528-101	Sequence 101, App
82	64.6	170	2	US-09-060-584-101	Sequence 101, App
83	64.6	170	2	US-09-413-140A-101	Sequence 101, App
84	64.6	197	2	US-09-248-796A-22808	Sequence 22808, A
85	64.6	231	2	US-10-104-047-2202	Sequence 2202, App
86	64.6	294	2	US-09-538-092-215	Sequence 215, App
87	64.6	310	2	US-09-902-540-15696	Sequence 15696, A
88	64.6	350	2	US-09-655-270A-17	Sequence 17, Appl
89	64.6	350	2	US-09-655-941-21	Sequence 21, Appl
90	64.6	350	2	US-09-955-597-21	Sequence 21, Appl
91	64.6	462	2	US-09-025-001A-16	Sequence 16, Appl
92	64.6	462	2	US-09-996-620-16	Sequence 16, Appl
93	64.6	521	2	US-09-026-001A-12	Sequence 12, Appl
94	64.6	521	2	US-09-996-620-12	Sequence 12, Appl
95	64.6	525	2	US-09-248-796A-15927	Sequence 15927, A
96	64.6	532	2	US-09-252-991A-17592	Sequence 17592, A
97	64.6	532	2	US-08-524-051-2	Sequence 2, Appl
98	64.6	554	1	US-08-524-051-2	Sequence 2, Appl
99	64.6	554	1	US-09-052-778-16	Sequence 16, Appl
100	64.6	554	2	US-09-052-778-16	Sequence 16, Appl

101	31	64.6	592	2	US-09-026-001A-14	Sequence 14, Appl1	174	30	62.5	466	2	US-09-134-000C-5176	Sequence 5176, Ap
102	31	64.6	592	2	US-09-996-620-14	Sequence 14, Appl1	175	30	62.5	469	2	US-09-107-532A-3672	Sequence 3672, Ap
103	31	64.6	613	2	US-09-026-001A-10	Sequence 10, Appl1	176	30	62.5	498	2	US-09-232-468A-18	Sequence 18, Appl1
104	31	64.6	613	2	US-09-996-620-10	Sequence 10, Appl1	177	30	62.5	498	2	US-08-686-968C-231	Sequence 231, Appl
105	31	64.6	617	2	US-09-252-991A-27878	Sequence 27878, A	178	30	62.5	498	2	US-09-784-984A-52	Sequence 52, Appl1
106	31	64.6	621	2	US-09-026-001A-18	Sequence 18, Appl1	179	30	62.5	499	2	US-09-328-352-5415	Sequence 5415, Ap
107	31	64.6	621	2	US-09-996-620-18	Sequence 18, Appl1	180	30	62.5	511	2	US-09-248-796A-16380	Sequence 16380, A
108	31	64.6	636	2	US-09-489-039A-8057	Sequence 8057, Ap	181	30	62.5	535	2	US-09-352-991A-22981	Sequence 22981, A
109	31	64.6	664	2	US-09-710-279-1258	Sequence 1258, Ap	182	30	62.5	545	2	US-09-270-767-46738	Sequence 46738, A
110	31	64.6	701	2	US-09-132-028-2	Sequence 2, Appl1	183	30	62.5	555	1	US-08-357-533A-9	Sequence 9, Appl1
111	31	64.6	723	2	US-09-564-805-339	Sequence 239, Appl	184	30	62.5	565	1	US-08-459-951-9	Sequence 9, Appl1
112	31	64.6	1098	2	US-10-104-047-2475	Sequence 2475, Ap	185	30	62.5	565	1	US-08-459-951-9	Sequence 9, Appl1
113	31	64.6	1103	2	US-09-162-373-1	Sequence 1, Appl1	186	30	62.5	567	1	US-08-361-873A-2	Sequence 2, Appl1
114	31	64.6	1103	2	US-09-467-946-1	Sequence 1, Appl1	187	30	62.5	567	1	US-08-483-926A-1	Sequence 1, Appl1
115	31	64.6	1123	2	US-09-949-016-9935	Sequence 9935, Ap	188	30	62.5	567	1	US-08-854-768-1	Sequence 1, Appl1
116	30	62.5	9	2	US-08-159-339A-320	Sequence 339, Appl	189	30	62.5	567	1	US-08-445-520B-9	Sequence 9, Appl1
117	30	62.5	9	2	US-08-159-339A-329	Sequence 339, Appl	190	30	62.5	567	1	US-08-737-045-1	Sequence 1, Appl1
118	30	62.5	10	2	US-08-159-339A-331	Sequence 331, Appl	191	30	62.5	567	1	US-08-451-946B-8	Sequence 8, Appl1
119	30	62.5	78	2	US-09-949-016-5824	Sequence 9824, Appl	192	30	62.5	567	1	US-08-446-938B-8	Sequence 8, Appl1
120	30	62.5	83	2	US-09-252-991A-27894	Sequence 27894, A	193	30	62.5	567	2	US-08-311-703A-8	Sequence 8, Appl1
121	30	62.5	89	2	US-09-543-681A-7711	Sequence 7711, Ap	194	30	62.5	567	2	US-08-446-939A-8	Sequence 8, Appl1
122	30	62.5	94	2	US-09-621-976-4330	Sequence 4330, Ap	195	30	62.5	567	2	US-09-183-543-8	Sequence 8, Appl1
123	30	62.5	108	2	US-09-328-352-6646	Sequence 4646, Ap	196	30	62.5	567	2	US-08-446-936A-8	Sequence 8, Appl1
124	30	62.5	117	2	US-09-540-336-3305	Sequence 3305, Ap	197	30	62.5	567	2	US-09-239-864A-11	Sequence 11, Appl1
125	30	62.5	122	2	US-09-949-016-8591	Sequence 8591, Ap	198	30	62.5	567	2	US-09-878-905-11	Sequence 11, Appl1
126	30	62.5	130	2	US-09-270-767-54956	Sequence 54956, A	199	30	62.5	567	2	US-09-267-963D-36	Sequence 36, Appl1
127	30	62.5	130	2	US-09-270-767-53071	Sequence 53071, A	200	30	62.5	567	4	PCT-US92-0932B-4	Sequence 4, Appl1
128	30	62.5	152	2	US-09-621-976-4924	Sequence 4924, Ap	201	30	62.5	604	2	US-09-605-703B-2158	Sequence 2158, Ap
129	30	62.5	154	2	US-09-621-976-4100	Sequence 4100, Ap	202	30	62.5	722	2	US-09-134-001C-5482	Sequence 5482, Ap
130	30	62.5	158	2	US-09-621-976-4282	Sequence 4282, Ap	203	30	62.5	841	2	US-09-902-540-14896	Sequence 14896, A
131	30	62.5	162	1	US-08-139-809-2	Sequence 2, Appl1	204	30	62.5	926	2	US-09-543-681A-6681	Sequence 6681, Ap
132	30	62.5	162	1	US-08-308-821A-2	Sequence 2, Appl1	205	30	62.5	1104	2	US-09-793-998-11	Sequence 11, Appl1
133	30	62.5	162	1	US-08-831-627-2	Sequence 2, Appl1	206	30	62.5	1194	2	US-09-092-508-2	Sequence 2, Appl1
134	30	62.5	162	2	US-09-495-491-2	Sequence 39739, A	207	30	62.5	1194	2	US-09-435-115-2	Sequence 26, Appl1
135	30	62.5	175	2	US-09-270-767-39739	Sequence 39739, A	208	30	62.5	1194	2	US-09-038-310-2	Sequence 2, Appl1
136	30	62.5	175	2	US-09-270-767-54956	Sequence 54956, A	209	30	62.5	1194	2	US-09-038-092-825	Sequence 825, Appl1
137	30	62.5	176	2	US-09-513-999C-5044	Sequence 5044, Ap	210	30	62.5	1194	2	US-09-949-016-6030	Sequence 6030, Ap
138	30	62.5	179	2	US-09-270-767-52208	Sequence 3208, A	211	30	62.5	1194	2	US-09-949-016-10065	Sequence 10065, A
139	30	62.5	179	2	US-09-270-767-47425	Sequence 47425, A	212	30	62.5	1196	2	US-09-949-016-10066	Sequence 10066, A
140	30	62.5	185	2	US-09-270-767-39277	Sequence 39277, A	213	30	62.5	1196	2	US-09-949-016-10066	Sequence 10066, A
141	30	62.5	185	2	US-09-270-767-54494	Sequence 54494, A	214	30	62.5	1205	2	US-09-092-508-16	Sequence 16, Appl1
142	30	62.5	187	2	US-09-248-796A-15530	Sequence 15530, A	215	30	62.5	1205	2	US-09-435-115-16	Sequence 16, Appl1
143	30	62.5	193	1	US-08-308-821A-6	Sequence 6, Appl1	216	30	62.5	1237	2	US-09-949-016-6842	Sequence 6842, Ap
144	30	62.5	193	2	US-08-831-627-6	Sequence 6, Appl1	217	30	62.5	1239	2	US-09-949-016-10063	Sequence 10063, A
145	30	62.5	193	2	US-08-765-381-15	Sequence 15, Appl1	218	30	62.5	1239	2	US-09-949-016-10064	Sequence 10064, A
146	30	62.5	193	2	US-09-495-491-6	Sequence 6, Appl1	219	30	62.5	1300	2	US-09-543-681A-4501	Sequence 4501, Ap
147	30	62.5	199	2	US-09-103-359-1	Sequence 11, Appl1	220	30	62.5	1979	2	US-09-949-016-6468	Sequence 6468, Ap
148	30	62.5	219	2	US-09-388-321B-21	Sequence 21, Appl1	221	30	62.5	2047	2	US-09-949-016-7404	Sequence 7404, Ap
149	30	62.5	221	2	US-09-248-796A-15073	Sequence 15073, A	222	30	62.5	2047	2	US-08-970-269A-29	Sequence 29, Appl1
150	30	62.5	226	2	US-09-198-452A-811	Sequence 811, Appl	223	30	62.5	2967	2	US-09-407-562-29	Sequence 29, Appl1
151	30	62.5	226	2	US-09-438-185A-763	Sequence 763, Appl	224	30	62.5	3959	1	US-08-970-269A-30	Sequence 30, Appl1
152	30	62.5	228	2	US-09-270-767-45453	Sequence 45453, A	225	30	62.5	3959	1	US-09-407-562-30	Sequence 30, Appl1
153	30	62.5	248	2	US-09-252-991A-20077	Sequence 20077, A	226	30	62.5	37	2	US-09-084-303B-283	Sequence 283, Appl
154	30	62.5	255	2	US-09-252-991A-19444	Sequence 19444, A	227	29	60.4	35	2	US-09-149-476-425	Sequence 425, Appl
155	30	62.5	277	2	US-09-538-092-710	Sequence 710, Appl	228	29	60.4	70	2	US-09-149-476-481	Sequence 481, Appl
156	30	62.5	308	2	US-10-014-269-31	Sequence 31, Appl1	229	29	60.4	72	1	US-08-542-363-13	Sequence 13, Appl1
157	30	62.5	308	2	US-10-002-974-31	Sequence 31, Appl1	230	29	60.4	72	1	US-09-100-087-13	Sequence 13, Appl1
158	30	62.5	342	2	US-09-584-568C-10	Sequence 10, Appl1	231	29	60.4	72	2	US-09-670-827-13	Sequence 13, Appl1
159	30	62.5	347	1	US-07-954-840A-2	Sequence 2, Appl1	232	29	60.4	82	2	US-09-827-949-13	Sequence 13, Appl1
160	30	62.5	379	2	US-09-248-796A-17132	Sequence 17132, A	233	29	60.4	80	2	US-09-084-303B-15	Sequence 15, Appl1
161	30	62.5	381	2	US-08-911-853-27	Sequence 27, Appl1	234	29	60.4	89	2	US-09-621-976-6758	Sequence 6768, Ap
162	30	62.5	381	2	US-09-479-409-27	Sequence 27, Appl1	235	29	60.4	89	2	US-09-446-938B-8	Sequence 8, Appl1
163	30	62.5	381	2	US-09-479-453-27	Sequence 27, Appl1	236	29	60.4	97	2	US-09-446-938B-8	Sequence 8, Appl1
164	30	62.5	391	2	US-08-968-563-11	Sequence 11, Appl1	237	29	60.4	97	2	US-09-311-021-64	Sequence 64, Appl1
165	30	62.5	391	2	US-08-969-683A-11	Sequence 11, Appl1	238	29	60.4	102	2	US-08-477-347-9	Sequence 9, Appl1
166	30	62.5	391	2	US-09-297-928-7	Sequence 7, Appl1	239	29	60.4	102	2	US-09-800-908-9	Sequence 9, Appl1
167	30	62.5	391	2	US-09-641-652-54	Sequence 54, Appl1	240	29	60.4	103	2	US-09-621-976-6448	Sequence 4448, Ap
168	30	62.5	391	2	US-09-308-207-11	Sequence 11, Appl1	241	29	60.4	109	2	US-09-270-767-61636	Sequence 61636, A
169	30	62.5	403	2	US-09-270-767-36436	Sequence 3436, A	242	29	60.4	115	2	US-09-583-110-2812	Sequence 2812, Ap
170	30	62.5	403	2	US-09-270-767-51653	Sequence 51653, A	243	29	60.4	115	2	US-09-107-433-3503	Sequence 3503, Appl
171	30	62.5	409	2	US-09-902-540-10657	Sequence 10657, A	244	29	60.4	127	1	US-08-162-146-3	Sequence 3, Appl1
172	30	62.5	410	2	US-09-489-039A-10689	Sequence 10689, A	245	29	60.4	127	2	US-09-314-127-3	Sequence 3, Appl1
173	30	62.5	442	2	US-09-540-236-2820	Sequence 2820, Ap	246	29	60.4	135	2	US-09-489-039A-10659	Sequence 10659, A

393	29	60.4	921	1	US-08-479-532-39	Sequence 39, Appl	466	28	58.3	60	2	US-09-583-110-3693	Sequence 3693, Ap
394	29	60.4	921	1	US-08-455-526-39	Sequence 39, Appl	467	28	58.3	74	2	US-09-122-135-3	Sequence 3, Appl1
395	29	60.4	921	1	US-08-455-525-39	Sequence 39, Appl	468	28	58.3	75	1	US-08-428-415-21	Sequence 21, Appl
396	29	60.4	921	2	US-09-139-491-39	Sequence 39, Appl	469	28	58.3	75	1	US-08-379-685-21	Sequence 21, Appl
397	29	60.4	921	2	US-09-754-250-5	Sequence 5, Appl1	470	28	58.3	75	1	US-08-854-029-21	Sequence 21, Appl
398	29	60.4	921	2	US-09-883-825-39	Sequence 39, Appl	471	28	58.3	75	2	US-08-428-762-21	Sequence 21, Appl
399	29	60.4	921	2	US-10-094-889-5	Sequence 5, Appl1	472	28	58.3	81	2	US-09-621-976-4507	Sequence 4507, Ap
400	29	60.4	921	4	PCT-US92-03222-39	Sequence 39, Appl	473	28	58.3	86	2	US-09-621-976-3953	Sequence 3953, Ap
401	29	60.4	921	1	US-07-872-644-45	Sequence 45, Appl	474	28	58.3	88	2	US-09-687-637B-45	Sequence 45, Appl
402	29	60.4	941	1	US-08-297-494-45	Sequence 45, Appl	475	28	58.3	88	2	US-09-270-767-58234	Sequence 58234, A
403	29	60.4	941	1	US-08-297-510-45	Sequence 45, Appl	476	28	58.3	88	2	US-09-270-767-60354	Sequence 60354, A
404	29	60.4	941	1	US-08-479-532-45	Sequence 45, Appl	477	28	58.3	102	1	US-08-808-982-8	Sequence 8, Appl1
405	29	60.4	941	1	US-08-455-526-45	Sequence 45, Appl	478	28	58.3	102	2	US-09-306-902A-8	Sequence 8, Appl1
406	29	60.4	941	1	US-08-455-525-45	Sequence 45, Appl	479	28	58.3	105	2	US-09-471-276-1565	Sequence 1565, Ap
407	29	60.4	941	2	US-09-139-491-45	Sequence 45, Appl	480	28	58.3	111	2	US-10-104-047-3831	Sequence 3831, Ap
408	29	60.4	941	2	US-09-883-825-45	Sequence 45, Appl	481	28	58.3	115	2	US-09-556-818-1	Sequence 1, Appl1
409	29	60.4	941	2	US-09-708-392-5	Sequence 5, Appl1	482	28	58.3	115	2	US-09-609-027B-6	Sequence 6, Appl1
410	29	60.4	941	2	US-09-949-016-6244	Sequence 6, Appl1	483	28	58.3	120	2	US-09-489-039K-12045	Sequence 12045, A
411	29	60.4	941	4	PCT-US92-03222-45	Sequence 45, Appl	484	28	58.3	121	2	US-09-180-864-2	Sequence 2, Appl1
412	29	60.4	942	1	US-07-872-644-43	Sequence 43, Appl	485	28	58.3	124	2	US-09-556-818-4	Sequence 4, Appl1
413	29	60.4	942	1	US-08-297-494-43	Sequence 43, Appl	486	28	58.3	124	2	US-09-556-818-5	Sequence 5, Appl1
414	29	60.4	942	1	US-08-297-510-43	Sequence 43, Appl	487	28	58.3	124	2	US-09-556-818-34	Sequence 34, Appl
415	29	60.4	942	1	US-08-479-532-43	Sequence 43, Appl	488	28	58.3	124	2	US-09-556-818-38	Sequence 38, Appl
416	29	60.4	942	1	US-08-455-526-43	Sequence 43, Appl	489	28	58.3	126	1	US-07-978-895-2	Sequence 2, Appl1
417	29	60.4	942	1	US-08-455-525-43	Sequence 43, Appl	490	28	58.3	126	1	US-08-473-119-2	Sequence 2, Appl1
418	29	60.4	942	2	US-09-139-491-43	Sequence 43, Appl	491	28	58.3	126	1	US-08-475-352-2	Sequence 2, Appl1
419	29	60.4	942	2	US-09-883-825-43	Sequence 43, Appl	492	28	58.3	126	2	US-09-170-699-2	Sequence 2, Appl1
420	29	60.4	942	4	PCT-US92-03222-43	Sequence 43, Appl	493	28	58.3	126	2	US-09-556-818-2	Sequence 2, Appl1
421	29	60.4	968	2	US-09-302-812-6	Sequence 6, Appl1	494	28	58.3	126	2	US-09-556-818-6	Sequence 6, Appl1
422	29	60.4	968	2	US-09-511-477-6	Sequence 6, Appl1	495	28	58.3	126	2	US-09-556-818-28	Sequence 28, Appl
423	29	60.4	968	2	US-09-511-507-6	Sequence 6, Appl1	496	28	58.3	126	6	5183884-2	Patent No. 5183884
424	29	60.4	968	2	US-09-973-451-6	Sequence 6, Appl1	497	28	58.3	129	2	US-10-104-047-2374	Sequence 2374, Ap
425	29	60.4	976	2	US-09-302-812-4	Sequence 4, Appl1	498	28	58.3	130	2	US-09-556-818-9	Sequence 9, Appl1
426	29	60.4	976	2	US-09-511-477-4	Sequence 4, Appl1	499	28	58.3	130	2	US-09-556-818-32	Sequence 32, Appl
427	29	60.4	976	2	US-09-511-507-4	Sequence 4, Appl1	500	28	58.3	130	2	US-09-556-818-42	Sequence 42, Appl
428	29	60.4	976	2	US-09-973-451-4	Sequence 4, Appl1	501	28	58.3	132	2	US-09-252-991A-21795	Sequence 21795, A
429	29	60.4	977	2	US-09-302-812-2	Sequence 2, Appl1	502	28	58.3	132	2	US-09-556-818-7	Sequence 7, Appl1
430	29	60.4	977	2	US-09-511-477-2	Sequence 2, Appl1	503	28	58.3	132	2	US-09-556-818-10	Sequence 10, Appl
431	29	60.4	977	2	US-09-511-507-2	Sequence 2, Appl1	504	28	58.3	132	2	US-09-556-818-4	Sequence 4, Appl
432	29	60.4	977	2	US-09-973-451-2	Sequence 2, Appl1	505	28	58.3	134	1	US-08-284-393B-13	Sequence 13, Appl
433	29	60.4	1238	2	US-09-252-991A-26363	Sequence 26363, A	506	28	58.3	134	2	US-08-759-628-9	Sequence 9, Appl1
434	29	60.4	1246	2	US-09-252-991A-23140	Sequence 23140, A	507	28	58.3	134	2	US-09-371-615A-7	Sequence 7, Appl1
435	29	60.4	1471	2	US-08-755-587-188	Sequence 188, App	508	28	58.3	134	2	US-09-462-941-12	Sequence 12, Appl
436	29	60.4	1589	2	US-08-755-587-189	Sequence 189, App	509	28	58.3	134	2	US-09-248-796A-26683	Sequence 26683, A
437	29	60.4	1620	1	US-08-542-363-2	Sequence 2, Appl1	510	28	58.3	134	4	PCT-US95-08950-13	Sequence 13, Appl
438	29	60.4	1620	2	US-09-100-089-2	Sequence 2, Appl1	511	28	58.3	134	6	5324640-2	Patent No. 5324640
439	29	60.4	1620	2	US-09-670-827-2	Sequence 2, Appl1	512	28	58.3	139	2	US-09-241-606-8	Sequence 8, Appl1
440	29	60.4	1620	2	US-09-827-949-2	Sequence 2, Appl1	513	28	58.3	141	2	US-09-270-767-33527	Sequence 33527, A
441	29	60.4	1711	2	US-08-369-822C-10	Sequence 10, Appl	514	28	58.3	141	2	US-09-270-767-33783	Sequence 33783, A
442	29	60.4	1711	2	US-08-582-776C-10	Sequence 10, Appl	515	28	58.3	141	2	US-09-270-767-48744	Sequence 48744, A
443	29	60.4	1712	2	US-08-434-831B-10	Sequence 10, Appl	516	28	58.3	141	2	US-09-270-767-49000	Sequence 49000, A
444	29	60.4	1752	2	US-09-949-002-294	Sequence 294, App	517	28	58.3	141	2	US-09-556-818-11	Sequence 11, Appl
445	29	60.4	1917	2	US-09-949-002-485	Sequence 485, App	518	28	58.3	147	2	US-09-556-818-44	Sequence 44, Appl
446	29	60.4	2183	1	US-08-348-891A-7	Sequence 7, Appl1	519	28	58.3	148	2	US-09-270-767-43240	Sequence 43240, A
447	29	60.4	2183	1	US-08-905-817-7	Sequence 7, Appl1	520	28	58.3	153	1	US-08-469-486-55	Sequence 55, Appl
448	29	60.4	3421	2	US-09-452-638-53	Sequence 53, Appl	521	28	58.3	153	1	US-08-469-658-55	Sequence 55, Appl
449	29	60.4	3421	2	US-09-121-587A-13	Sequence 13, Appl	522	28	58.3	172	1	US-08-438-153B-2	Sequence 2, Appl1
450	28.5	59.4	149	2	US-09-270-767-3323	Sequence 3323, A	523	28	58.3	172	1	US-08-443-883A-2	Sequence 2, Appl1
451	28.5	59.4	149	2	US-09-270-767-48540	Sequence 48540, A	524	28	58.3	172	1	US-08-631-328-2	Sequence 2, Appl1
452	28	58.3	15	2	US-09-149-476-510	Sequence 510, App	525	28	58.3	172	1	US-08-455-524B-2	Sequence 2, Appl1
453	28	58.3	21	1	US-08-335-888A-15	Sequence 15, Appl	526	28	58.3	172	1	US-08-455-021B-2	Sequence 2, Appl1
454	28	58.3	21	1	US-08-693-697-15	Sequence 15, Appl	527	28	58.3	172	2	US-09-045-467-2	Sequence 2, Appl1
455	28	58.3	21	1	US-08-693-697-34	Sequence 34, Appl	528	28	58.3	172	2	US-08-554-395A-18	Sequence 18, Appl
456	28	58.3	21	1	US-08-588-526-4	Sequence 4, Appl1	529	28	58.3	172	2	US-08-616-904-2	Sequence 2, Appl1
457	28	58.3	21	2	US-08-693-696-15	Sequence 15, Appl	530	28	58.3	172	2	US-09-599-413-2	Sequence 2, Appl1
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596	28	58.3	325	1	US-08-087-797-2	Sequence 2, Appl	669	28	58.3	631	2	US-08-927-219-127	Sequence 109, App
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887	27	56.2	222	2	US-09-465-901-34	Sequence 34, Appli	960	27	56.2	346	2	US-09-902-540-15109	Sequence 8, Appli
888	27	56.2	224	2	US-09-270-767-43554	Sequence 43554, A	961	27	56.2	357	2	US-09-535-909-8	Sequence 30177, A
889	27	56.2	228	2	US-09-336-536-4	Sequence 4, Appli	962	27	56.2	357	2	US-09-252-991A-26177	Sequence 26177, A
890	27	56.2	228	2	US-09-336-536-11	Sequence 11, Appli	963	27	56.2	357	2	US-10-104-047-3031	Sequence 9473, Ap
891	27	56.2	235	2	US-10-104-047-3131	Sequence 3131, Ap	964	27	56.2	362	2	US-09-489-039A-9473	Sequence 9473, Ap
892	27	56.2	236	2	US-09-634-137-4	Sequence 4, Appli	965	27	56.2	362	2	US-09-489-039A-9473	Sequence 32637, A
893	27	56.2	237	2	US-08-679-493A-73	Sequence 73, Appli	966	27	56.2	374	4	US-09-252-991A-32637	Sequence 7, Appli
894	27	56.2	237	2	US-09-107-513A-4963	Sequence 4963, Ap	967	27	56.2	382	1	US-08-470-299-7	Sequence 10, Appli
895	27	56.2	243	2	US-09-188-930-295	Sequence 295, App	968	27	56.2	382	1	US-09-248-796A-18423	Sequence 18423, A
896	27	56.2	243	2	US-09-140-804-2	Sequence 2, Appli	969	27	56.2	386	1	US-08-559-803B-75	Sequence 75, Appli
897	27	56.2	243	2	US-09-336-536-3	Sequence 3, Appli	970	27	56.2	386	2	US-09-175-828-75	Sequence 75, Appli
898	27	56.2	243	2	US-09-336-536-10	Sequence 10, Appli	971	27	56.2	386	2	US-09-328-352-1608	Sequence 7608, Ap
899	27	56.2	243	2	US-09-686-838B-2	Sequence 2, Appli	972	27	56.2	386	2	US-09-753-143-75	Sequence 4, Appli
900	27	56.2	243	2	US-09-312-283C-295	Sequence 295, App	973	27	56.2	387	1	US-08-470-299-4	Sequence 4, Appli
901	27	56.2	243	2	US-09-866-028-42	Sequence 42, Appli	974	27	56.2	392	2	US-09-252-991A-20053	Sequence 20053, A
902	27	56.2	243	2	US-09-944-457-42	Sequence 42, Appli	975	27	56.2	393	2	US-09-248-796A-15182	Sequence 15182, A
903	27	56.2	243	2	US-09-945-584-42	Sequence 42, Appli	976	27	56.2	393	2	US-09-248-796A-15182	Sequence 15182, A

977 27 56.2 401 1 US-08-805-118-1 Sequence 1, Appli
978 27 56.2 401 2 US-09-391-958-1 Sequence 1, Appli
979 27 56.2 401 2 US-09-248-796A-15172 Sequence 15172, A
980 27 56.2 401 2 US-10-272-490-38 Sequence 38, Appl
981 27 56.2 402 2 US-10-272-490-10 Sequence 10, Appl
982 27 56.2 405 2 US-09-489-039A-9126 Sequence 9126, Ap
983 27 56.2 406 2 US-08-934-494-2 Sequence 2, Appli
984 27 56.2 406 2 US-09-143-068-2 Sequence 2, Appli
985 27 56.2 406 2 US-09-143-707-2 Sequence 2, Appli
986 27 56.2 406 2 US-09-202-089-2 Sequence 2, Appli
987 27 56.2 406 2 US-09-511-133-2 Sequence 2, Appli
988 27 56.2 406 2 US-09-690-169-2 Sequence 2, Appli
989 27 56.2 406 2 US-09-511-631-2 Sequence 2, Appli
990 27 56.2 406 2 US-09-461-325-206 Sequence 206, App
991 27 56.2 406 2 US-09-690-189-2 Sequence 2, Appli
992 27 56.2 406 2 US-10-012-542-206 Sequence 206, App
993 27 56.2 406 2 US-10-115-123-206 Sequence 206, App
994 27 56.2 407 2 US-09-252-991A-32423 Sequence 32423, A
995 27 56.2 408 2 US-10-272-490-56 Sequence 36, Appl
996 27 56.2 410 2 US-09-328-352-7301 Sequence 7301, Ap
997 27 56.2 410 2 US-10-272-490-24 Sequence 24, Appl
998 27 56.2 412 2 US-09-248-796A-22404 Sequence 22404, A
999 27 56.2 413 2 US-09-585-173B-44 Sequence 44, Appl
1000 27 56.2 425 2 US-09-538-092-983 Sequence 983, App

ALIGNMENTS

RESULT 1
US-08-787-547-101
Sequence 101, Application US/08787547
Patent No. 5783567
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Curley, Joanne M.
APPLICANT: Langer, Robert S.
TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY
TITLE OF INVENTION: OF NUCLEIC ACID
NUMBER OF SEQUENCES: 107
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/787, 547
FILING DATE: 22-JAN-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08191/003001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 101:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide

US-08-787-547-101

Query Match 100.0%; Score 48; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 4,6e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 1 KLPOLCTEL 9

RESULT 2

US-09-601-729-274

Sequence 274, Application US/09601729

Patent No. 6683052

GENERAL INFORMATION:

APPLICANT: THIAM, KADER

APPLICANT: AURIAULT, CLAUDE

APPLICANT: GRAS-MASSE, HELENE

APPLICANT: LOING, ESTELLE

APPLICANT: VERMARDE, CLAUDE

APPLICANT: GUILLET, JEAN GERARD

TITLE OF INVENTION: LIPOPEPTIDES CONTAINING AN INTERFERON FRAGMENT AND USES

TITLE OF INVENTION: THEREOF IN PHARMACEUTICAL COMPOSITIONS

FILE REFERENCE: USB-97-RU-IN

CURRENT APPLICATION NUMBER: US/09/601,729

CURRENT FILING DATE: 2000-11-20

PRIOR APPLICATION NUMBER: PCT/FR99/00259

PRIOR FILING DATE: 1999-02-05

PRIOR APPLICATION NUMBER: 98 01439

PRIOR FILING DATE: 1998-02-06

NUMBER OF SEQ ID NOS: 281

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 274

LENGTH: 9

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE: Description of Artificial Sequence: Synthetic

OTHER INFORMATION: peptide

US-09-601-729-274

Query Match 100.0%; Score 48; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4,6e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 1 KLPOLCTEL 9

RESULT 3

US-08-934-915-159

Sequence 159, Application US/08934915

Patent No. 5932412

GENERAL INFORMATION:

APPLICANT: DILLNER, JOAKIM

APPLICANT: DILLNER, LENA

APPLICANT: CHENG, HWEI-MING

TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN

TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,

TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,

TITLE OF INVENTION: USEFUL IN IMMUNOSSAY FOR

NUMBER OF SEQUENCES: 193

CORRESPONDENCE ADDRESS:

ADDRESSEE: MASON & ASSOCIATES, P.A.

STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500

CITY: CLEARWATER

STATE: FLORIDA

COUNTRY: U.S.A.

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible
OPERATING SYSTEM: Windows 3.0
SOFTWARE: Microsoft Word 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/934,915
FILING DATE: 22-SEP-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/949,836
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: LOUISE A. Foutch
REGISTRATION NUMBER: 37,133
REFERENCE/DOCKET NUMBER: 1946.6
TELECOMMUNICATION INFORMATION:
TELEPHONE: 813-538-3800
TELEFAX: 813-538-3820
TELEX:
INFORMATION FOR SEQ ID NO: 159:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-934-915-159

Query Match 100.0%; Score 48; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.029;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPPOLCTEL 9
|||
2 KLPPOLCTEL 10

Db

RESULT 4
US-08-363-586-4
Sequence 4, Application US/08363586
Patent No. 5629161
GENERAL INFORMATION:
APPLICANT: Mueller, Martin
APPLICANT: Gissmann, Lutz
TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
TITLE OF INVENTION: Peptides for the Diagnostic Purpose
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESS: Dunner
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,586
FILING DATE: 23-DEC-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/909,296
FILING DATE: 09-JUL-1992
APPLICATION NUMBER: EP 91111720.8
FILING DATE: 13-JUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Madler, Linda A.
REGISTRATION NUMBER: 33,218
REFERENCE/DOCKET NUMBER: 02481-1195-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000

TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-363-586-4

Query Match 100.0%; Score 48; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.043;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPPOLCTEL 9
|||
11 KLPPOLCTEL 19

Db

RESULT 5
US-09-980-523A-4
Sequence 4, Application US/09980523A
Patent No. 6783763
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCES
APPLICANT: FERRIER, ESTELLE
TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
TITLE OF INVENTION: PARTICULARLY IN VACCINATION
FILE REFERENCE: WOBI AO INS
CURRENT APPLICATION NUMBER: US/09/980,523A
CURRENT FILING DATE: 2002-04-29
PRIOR APPLICATION NUMBER: PCT/FR00/01513
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: FR 99/07012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 4
LENGTH: 30
TYPE: PPT
ORGANISM: Human Papillomavirus
US-09-980-523A-4

Query Match 100.0%; Score 48; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.043;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPPOLCTEL 9
|||
4 KLPPOLCTEL 12

Db

RESULT 6
US-09-701-080C-18
Sequence 18, Application US/09701080C
Patent No. 6664054
GENERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REFERENCE: N73477C GCM
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR FILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: GB 9900157.0
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 18

LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match 100.0%; Score 48; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 11 KLPOLCTEL 19

RESULT 7
US-09-980-523A-2
Sequence 2, Application US/09980523A
Patent No. 6783763
GENERAL INFORMATION:
APPLICANT: CHOIPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCES
APPLICANT: FERRIS, ESTELLE
TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
FILE REFERENCE: MOBI AO INS
CURRENT APPLICATION NUMBER: US/09/980,523A
CURRENT FILING DATE: 2002-04-29
PRIOR APPLICATION NUMBER: PCT/FR00/01513
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: FR 99/07012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match 100.0%; Score 48; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.22;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26

RESULT 8
US-08-316-239B-3
Sequence 3, Application US/08316239B
Patent No. 5679509
GENERAL INFORMATION:
APPLICANT: Wheeler, Cosette M.
APPLICANT: Parmenter, Cheryl A.
TITLE OF INVENTION: Methods and a Diagnostic Aid for
TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
TITLE OF INVENTION: Cervical Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Jagtiani & Associates
STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtiani, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-3

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26

RESULT 9
US-08-316-239B-4
Sequence 4, Application US/08316239B
Patent No. 5679509
GENERAL INFORMATION:
APPLICANT: Wheeler, Cosette M.
APPLICANT: Parmenter, Cheryl A.
TITLE OF INVENTION: Methods and a Diagnostic Aid for
TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
TITLE OF INVENTION: Cervical Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Jagtiani & Associates
STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtiani, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant

MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-2398-4

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26

RESULT 10
US-08-860-165-14
Sequence 14, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860.165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentln Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14

Query Match 100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.24;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 87 KLPOLCTEL 95

RESULT 11
US-09-359-382-14
Sequence 14, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359.382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860.165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentln Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT

ORGANISM: Human papillomavirus type 16
US-09-359-382-14

Query Match 100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.24;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 87 KLPOLCTEL 95

RESULT 12
US-08-117-083-10
Sequence 10, Application US/08117083
Patent No. 5719054
GENERAL INFORMATION:
APPLICANT: Boursnell, Michael E.
APPLICANT: Inglis, Stephen C.
APPLICANT: Munro, Alan J.
TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSEE: Walter H. Dregger
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentln Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117.083
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dregger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-58783
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 182 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..182
OTHER INFORMATION:
OTHER INFORMATION: /note= "Xaa refers to stop codon in
US-08-117-083-10

Query Match 100.0%; Score 48; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.26;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 19 KLPOLCTEL 27

RESULT 13
US-09-462-993-1
Sequence 1, Application US/09462993

```
; Patent No. 6884786
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOUL, Jean-Marc
; APPLICANT: BIZOUARN, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone B6*TMF.
US-09-462-993-1
```

```
Query Match          100.0%; Score 48; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.34;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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QY 1 KLPOLCTEL 9
Db 46 KLPOLCTEL 54
```

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RESULT 14
US-08-860-165-10
; Sequence 10, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; PRIOR FILING DATE: 1997-09-22
; PRIOR APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10
```

```
Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.37;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26
```

```
RESULT 15
US-09-359-382-10
```

```
; Sequence 10, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; PRIOR FILING DATE: 1999-07-23
; PRIOR APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-359-382-10
```

```
Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.37;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26
```

```
RESULT 16
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Patent No. 6428807
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; PRIOR FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1
```

```
Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.37;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26
```

```
RESULT 17
US-09-485-885-4
; Sequence 4, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: BRUCK, Claudine
; APPLICANT: CABEZON Silva, Teresa
```

```
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4
```

```
Query Match          100.0%; Score 48; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.38;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
Db 124 KLPOLCTEL 132
```

```
RESULT 18
US-09-485-885-10
; Sequence 10, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
```

```
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-10
```

```
Query Match          100.0%; Score 48; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.41;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
Db 143 KLPOLCTEL 151
```

```
RESULT 19
US-09-485-885-6
; Sequence 6, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
```

```
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
```

```
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-6
```

```
Query Match          100.0%; Score 48; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.52;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
Db 124 KLPOLCTEL 132
```

```
RESULT 20
US-09-485-885-14
; Sequence 14, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
```

```
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-14
```

```
Query Match          100.0%; Score 48; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.54;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
Db 143 KLPOLCTEL 151
```

```
RESULT 21
US-08-466-285-2
; Sequence 2, Application US/08466285
; Patent No. 5753233
; GENERAL INFORMATION:
```

```
APPLICANT: Bleul, Conrad
APPLICANT: Gissmann, Lutz
APPLICANT: Muller, Martin
TITLE OF INVENTION: Seroreactive Epitopes On Proteins Of
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSER: Finnegan, Henderson, Farabow, Garrett &
```

ADDRESS: Dunner
STREET: 1300 I Street, N.W., Suite 700
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,285
FILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/947,992
FILING DATE: 21-SEP-1992
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/696,953
FILING DATE: 08-MAY-1991
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: P 40 15 044.5
FILING DATE: 10-MAY-1990
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Manspeizer, David A.
REGISTRATION NUMBER: 37,540
REFERENCE/DOCKET NUMBER: 05552.1075-03000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-466-285-2

Query Match 89.6%; Score 43; DB 1; Length 32;
Best Local Similarity 88.9%; Pred. No. 0.37;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPDLCTEL 9
DB 8 KLPDLCTEL 16

RESULT 22
US-08-164-768-2
Sequence 2, Application US/08164768
Patent No. 6322794
GENERAL INFORMATION:
APPLICANT: BLEUL, Conrad
APPLICANT: GISSMANN, Lutz
APPLICANT: MULLER, Martin
TITLE OF INVENTION: SEROREACTIVE EPITOPES ON PROTEINS OF
TITLE OF INVENTION: HUMAN PAPILLOMA VIRUS (HPV) 18
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: FINNEGAN, HENDERSON, PARABOW, GARRETT &
ADDRESSEE: DUNNER, L.L.P.
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: DC

COUNTRY: USA
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Forman, David S.
REGISTRATION NUMBER: 33,694
REFERENCE/DOCKET NUMBER: 05552.1075-02000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-164-768-2

Query Match 89.6%; Score 43; DB 2; Length 32;
Best Local Similarity 88.9%; Pred. No. 0.37;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPDLCTEL 9
DB 8 KLPDLCTEL 16

RESULT 23
US-08-247-904B-10
Sequence 10, Application US/08247904B
Patent No. 5981699
GENERAL INFORMATION:
APPLICANT: Rolfe, Mark
APPLICANT: Eckstein, Jens W.
TITLE OF INVENTION: Human Ubiquitin Conjugating Enzyme
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley, Hoag & Elliot
STREET: One Post Office Square
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII(text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/247,904B
FILING DATE: 23-MAY-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: MIV-029.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 832-1000
TELEFAX: (617) 832-7000
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 amino acids
TYPE: amino acid

```

; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-247-904B-10

Query Match      89.6%; Score 43; DB 1; Length 158;
Best Local Similarity 88.9%; Pred. No. 1.8;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KLPOLCTEL 9
      ||| ||| |||
Db      13 KLPDLCTEL 21

RESULT 24
US-08-767-942A-19
; Sequence 19, Application US/08767942A
; Patent No. 6068982
; GENERAL INFORMATION:
; APPLICANT: Rolfe, Mark
; APPLICANT: Chiu, M. Isabel
; APPLICANT: Berlin, Vivian
; APPLICANT: Damagnez, Veronique
; APPLICANT: Draetta, Giulio
; APPLICANT: Guillaume, Cottarel
; TITLE OF INVENTION: UBIQUITIN CONJUGATING ENZYMES
; NUMBER OF SEQUENCES: 45
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: FOLEY, HOAG & ELIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02108-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/767,942A
; FILING DATE: 17-DEC-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Vincent, Matthew P.
; REGISTRATION NUMBER: 36,709
; REFERENCE/DOCKET NUMBER: MIV-029.04
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 19:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 158 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-767-942A-19

Query Match      89.6%; Score 43; DB 2; Length 158;
Best Local Similarity 88.9%; Pred. No. 1.8;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KLPOLCTEL 9
      ||| ||| |||
Db      13 KLPDLCTEL 21

RESULT 25
US-08-117-083-14
; Sequence 14, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Bourtenell, Michael E.
; APPLICANT: Ingils, Stephen C.
; APPLICANT: Munro, Alan J.
```

```

; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; TITLE OF INVENTION: Papilloma Virus Proteins
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Walter H. Dregger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dregger, Walter H.
; REGISTRATION NUMBER: 24,190
; REFERENCE/DOCKET NUMBER: A-58783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 271 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..271
; OTHER INFORMATION: /note= "Xaa refers to stop codon in
; OTHER INFORMATION: the open reading frame."
US-08-117-083-14

Query Match      89.6%; Score 43; DB 1; Length 271;
Best Local Similarity 88.9%; Pred. No. 3.1;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      1 KLPOLCTEL 9
      ||| ||| |||
Db      14 KLPDLCTEL 22

RESULT 26
US-09-485-885-21
; Sequence 21, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 21
; LENGTH: 278
; TYPE: PRT
```

; ORGANISM: Homo sapien
US-09-485-885-21

Query Match 89.6%; Score 43; DB 2; Length 278;
Best Local Similarity 88.9%; Pred. No. 3.2;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 124 KLPDLCTEL 132

RESULT 27
US-09-485-885-23
; Sequence 23, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisee, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FaastSeq for Windows Version 3.0
; SEQ ID NO 23
; LENGTH: 383
; TYPE: PRF
; ORGANISM: Homo sapien
US-09-485-885-23

Query Match 89.6%; Score 43; DB 2; Length 383;
Best Local Similarity 88.9%; Pred. No. 4.3;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 124 KLPDLCTEL 132

RESULT 28
US-09-543-681A-8111
; Sequence 811, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETTON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; CURRENT FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO 8111
; LENGTH: 211
; TYPE: PRF
; ORGANISM: Proteus mirabilis
US-09-543-681A-8111

Query Match 72.9%; Score 35; DB 2; Length 211;
Best Local Similarity 77.8%; Pred. No. 68;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 124 KLPDLCTEL 132

Db 187 KFNOLCTEL 195

RESULT 29
US-07-909-122-2
; Sequence 2, Application US/07909122
; Patent No. 5415995
; GENERAL INFORMATION:
; APPLICANT: SCHOOLNIK, GARY K.
; APPLICANT: PALEFSKY, JOEL M.
; TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
; TITLE OF INVENTION: VIRUS
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 Page Mill Road
; CITY: Palo Alto
; STATE: California
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/909,122
; FILING DATE: 19920706
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: BENZ, WILLIAM H.
; REGISTRATION NUMBER: 25,952
; REFERENCE/DOCKET NUMBER: 28600-20105.01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 813-5600
; TELEFAX: (415) 494-0792
; TELEX: 706141
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 15 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
US-07-909-122-2

Query Match 70.8%; Score 34; DB 1; Length 15;
Best Local Similarity 100.0%; Pred. No. 7.5;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLC 6
Db 10 KLPOLC 15

RESULT 30
US-08-075-541D-52
; Sequence 52, Application US/0807541D
; Patent No. 6183745
; GENERAL INFORMATION:
; APPLICANT: TINDLE, ROBERT
; APPLICANT: FERNANDO, GERMAIN
; APPLICANT: FRAZER, IAN
; TITLE OF INVENTION: SUBUNIT PAPILLOMA VIRUS VACCINE AND
; TITLE OF INVENTION: PEPTIDES FOR USE THEREIN
; NUMBER OF SEQUENCES: 56
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: PANITCH SCHWARZE JACOBS & NADEL, P.C.
; STREET: 1601 MARKET STREET, 36TH FLOOR
; CITY: PHILADELPHIA
; STATE: PENNSYLVANIA
; COUNTRY: USA
; ZIP: 19103-2398
; COMPUTER READABLE FORM:

MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/075,541D
FILING DATE: 10-JUN-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU pk 3876
FILING DATE: 12-DEC-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: pcc/aus1/00575
FILING DATE: 12-DEC-1991
ATTORNEY/AGENT INFORMATION:
NAME: NADEL, ALAN S
REGISTRATION NUMBER: 27,363
REFERENCE/DOCKET NUMBER: 8795-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-567-2020
TELEFAX: 215-567-2991
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-075-541D-52

Query Match 70.8%; Score 34; DB 2; Length 15;
Best Local Similarity 100.0%; Pred. No. 7.5;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLC 6
Db 10 KLPOLC 15

RESULT 31
US-08-363-586-3
Sequence 3, Application US/08363586
Patent No. 5629161
GENERAL INFORMATION:
APPLICANT: Mueller, Martin
APPLICANT: Gissmann, Lutz
TITLE OF INVENTION: Use of HPV-16 B6 and E7-Gene Derived
TITLE OF INVENTION: Peptides for the Diagnostic Purpose
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESSEE: Dunner
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,586
FILING DATE: 23-DEC-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/909,296
FILING DATE: 09-JUL-1992
APPLICATION NUMBER: BP 91111720.8
FILING DATE: 13-JUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Madler, Linda A.

REGISTRATION NUMBER: 33,218
REFERENCE/DOCKET NUMBER: 02481-1195-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 23 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-363-586-3

Query Match 70.8%; Score 34; DB 1; Length 23;
Best Local Similarity 100.0%; Pred. No. 12;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLC 6
Db 18 KLPOLC 23

RESULT 32
US-09-673-395A-248
Sequence 248, Application US/09673395A
Patent No. 6620923
GENERAL INFORMATION:
APPLICANT: SPECHT, THOMAS
APPLICANT: HINZMANN, BERND
APPLICANT: SCHMITT, ARMIN
APPLICANT: PILARSKY, CHRISTIAN
APPLICANT: DAHL, EDGAR
APPLICANT: ROSENTHAL, ANDRE
TITLE OF INVENTION: HUMAN NUCLEIC ACID SEQUENCES FROM UTERUS TUMOR TISSUE
FILE REFERENCE: ALBRE-12
CURRENT APPLICATION NUMBER: US/09/673,395A
CURRENT FILING DATE: 2000-10-17
NUMBER OF SEQ ID NOS: 637
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 248
LENGTH: 161
TYPE: PRT
ORGANISM: Homo sapiens
US-09-673-395A-248

Query Match 70.8%; Score 34; DB 2; Length 161;
Best Local Similarity 85.7%; Pred. No. 79;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCT 7
Db 54 KLPOLCT 60

RESULT 33
US-08-481-968A-11
Sequence 11, Application US/08481968A
Patent No. 6300490
GENERAL INFORMATION:
APPLICANT: Huber, Brian
APPLICANT: Richards, Cynthia
TITLE OF INVENTION: Molecular Constructs Comprising a Carcinoembryonic Antigen (C
TITLE OF INVENTION: Transcriptional Regulatory Region
FILE REFERENCE: PB1087US4
CURRENT APPLICATION NUMBER: US/08/481,968A
CURRENT FILING DATE: 1998-06-07
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin version 3.0
SEQ ID NO 11
LENGTH: 341
TYPE: PRT
ORGANISM: Varicella zoster
US-08-481-968A-11

Query Match 70.8%; Score 34; DB 2; Length 341;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTE 8
|:|:|:|:
Db 262 KLPBLCGE 269

RESULT 34
US-08-154-712B-11
; Sequence 11, Application US/08154712B
; Patent No. 6337209
; GENERAL INFORMATION:
; APPLICANT: Huber, Brian
; APPLICANT: Richards, Cynthia
; TITLE OF INVENTION: Molecular Constructs Containing a Carcinoembryonic Antigen Regu
; FILE REFERENCE: PB1087US3
; CURRENT APPLICATION NUMBER: US/08/154,712B
; CURRENT FILING DATE: 1993-11-19
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 11
; LENGTH: 341
; TYPE: PRT
; ORGANISM: Varicella zoster
US-08-154-712B-11

Query Match 70.8%; Score 34; DB 2; Length 341;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTE 8
|:|:|:|:
Db 262 KLPBLCGE 269

RESULT 35
US-09-947-925A-11
; Sequence 11, Application US/09947925A
; Patent No. 6696960
; GENERAL INFORMATION:
; APPLICANT: Huber, Brian
; APPLICANT: Richards, Cynthia
; TITLE OF INVENTION: Molecular Constructs Containing a Carcinoembryonic
; FILE REFERENCE: PB1087US3
; CURRENT APPLICATION NUMBER: US/09/947,925A
; CURRENT FILING DATE: 2001-09-06
; PRIOR APPLICATION NUMBER: US/08/154,712
; PRIOR FILING DATE: 1993-11-19
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: Patentin version 3.0
; SEQ ID NO 11
; LENGTH: 341
; TYPE: PRT
; ORGANISM: Varicella zoster
US-09-947-925A-11

Query Match 70.8%; Score 34; DB 2; Length 341;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTE 8
|:|:|:|:
Db 262 KLPBLCGE 269

RESULT 36
US-09-198-452A-1036

; Sequence 1036, Application US/09198452A
; Patent No. 6559294
; GENERAL INFORMATION:
; APPLICANT: Griffiths, R.
; TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragment
; TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, prev
; TITLE OF INVENTION: and treatment of infection
; FILE REFERENCE: 9710-003-999
; CURRENT APPLICATION NUMBER: US/09/198,452A
; CURRENT FILING DATE: 1998-11-24
; NUMBER OF SEQ ID NOS: 6849
; SEQ ID NO 1036
; LENGTH: 504
; TYPE: PRT
; ORGANISM: Chlamydia pneumoniae
; FEATURE:
; NAME/KEY: SITE
; LOCATION: 1...504
; OTHER INFORMATION: Xaa=unknown or other
US-09-198-452A-1036

Query Match 70.8%; Score 34; DB 2; Length 504;
Best Local Similarity 55.6%; Pred. No. 2.5e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
|:|:|:|:
Db 206 KIPDLCSQL 214

RESULT 37
US-09-438-185A-966
; Sequence 966, Application US/09438185A
; Patent No. 6822071
; GENERAL INFORMATION:
; APPLICANT: Stephens, Richard
; APPLICANT: Mitchell, Wayne
; APPLICANT: Kaiman, Sue
; APPLICANT: Davis, Ronald
; APPLICANT: The Regents of the University of California
; TITLE OF INVENTION: Chlamydia Pneumoniae Genome Sequence
; FILE REFERENCE: 018941-00041US
; CURRENT APPLICATION NUMBER: US/09/438,185A
; CURRENT FILING DATE: 2002-03-13
; PRIOR APPLICATION NUMBER: US 60/108,279
; PRIOR FILING DATE: 1998-11-12
; PRIOR APPLICATION NUMBER: US 60/128,606
; PRIOR FILING DATE: 1999-04-08
; NUMBER OF SEQ ID NOS: 1074
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 966
; LENGTH: 504
; TYPE: PRT
; ORGANISM: Chlamydia pneumoniae
; FEATURE:
; OTHER INFORMATION: CPN0964
US-09-438-185A-966

Query Match 70.8%; Score 34; DB 2; Length 504;
Best Local Similarity 55.6%; Pred. No. 2.5e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
|:|:|:|:
Db 206 KIPDLCSQL 214

RESULT 38
US-09-949-016-10663
; Sequence 10663, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.

```

; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: C0001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10663
; LENGTH: 522
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-10663
```

```
Query Match      70.8%; Score 34; DB 2; Length 522;
Best Local Similarity 66.7%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;
```

```
QY      1 KLPOCTEL 9
Db      177 KLPOFCALH 185
```

```

RESULT 39
US-09-747-259-18
; Sequence 18, Application US/09747259
; Patent No. 6569645
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Chen, Jian
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul
; APPLICANT: Grimaldi, Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Li, Hanzhong
; APPLICANT: Hillan, Kenneth
; APPLICANT: Tumas, Daniel
; APPLICANT: Vanlookeren, Menno
; APPLICANT: Vandlen, Richard
; APPLICANT: Watanabe, Colin
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William
; APPLICANT: Yansura, Daniel
; TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
; FILE REFERENCE: P1381R1C1P1 (US)
; CURRENT APPLICATION NUMBER: US/09/747,259
; CURRENT FILING DATE: 2000-12-20
; PRIOR APPLICATION NUMBER: US 09/311,832
; PRIOR FILING DATE: 1999-05-14
; PRIOR APPLICATION NUMBER: US 60/172,096
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: PCT/US99/31274
; PRIOR FILING DATE: 1999-12-30
; PRIOR APPLICATION NUMBER: US 60/175,481
; PRIOR FILING DATE: 2000-01-11
; PRIOR APPLICATION NUMBER: PCT/US00/04341
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/US00/05841
; PRIOR FILING DATE: 2000-03-02
; PRIOR APPLICATION NUMBER: US 60/191,007
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: PCT/US00/07532
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: PCT/US00/15264
; PRIOR FILING DATE: 2000-06-02
; PRIOR APPLICATION NUMBER: US 60/213,087
```

```

; PRIOR FILING DATE: 2000-06-22
; PRIOR APPLICATION NUMBER: US 09/644,848
; PRIOR FILING DATE: 2000-08-22
; PRIOR APPLICATION NUMBER: PCT/US00/23328
; PRIOR FILING DATE: 2000-08-24
; PRIOR APPLICATION NUMBER: US 60/242,837
; PRIOR FILING DATE: 2000-10-24
; PRIOR APPLICATION NUMBER: PCT/US00/30873
; PRIOR FILING DATE: 2000-11-10
; PRIOR APPLICATION NUMBER: US 60/253,646
; PRIOR FILING DATE: 2000-11-28
; PRIOR APPLICATION NUMBER: PCT/US00/32678
; PRIOR FILING DATE: 2000-12-01
; NUMBER OF SEQ ID NOS: 39
; SEQ ID NO 18
; LENGTH: 728
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-747-259-18
```

```
Query Match      70.8%; Score 34; DB 2; Length 728;
Best Local Similarity 75.0%; Pred. No. 3.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      2 LPOLCTEL 9
Db      493 LPOLCSHL 500
```

```

RESULT 40
US-09-816-744-18
; Sequence 18, Application US/09816744
; Patent No. 6579520
; GENERAL INFORMATION:
; APPLICANT: Chen, Jian
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Fong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul
; APPLICANT: Grimaldi, Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Li, Hanzhong
; APPLICANT: Hillan, Kenneth
; APPLICANT: Tumas, Daniel
; APPLICANT: Vanlookeren, Menno
; APPLICANT: Vandlen, Richard
; APPLICANT: Watanabe, Colin
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William
; APPLICANT: Yansura, Daniel
; TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
; FILE REFERENCE: P1381R1C1P2 (US)
; CURRENT APPLICATION NUMBER: US/09/816,744
; CURRENT FILING DATE: 2001-03-22
; Prior application data removed - consult PAM or file wrapper
; NUMBER OF SEQ ID NOS: 39
; SEQ ID NO 18
; LENGTH: 728
; TYPE: PRT
; ORGANISM: Homo Sapien
US-09-816-744-18
```

```
Query Match      70.8%; Score 34; DB 2; Length 728;
Best Local Similarity 75.0%; Pred. No. 3.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      2 LPOLCTEL 9
Db      493 LPOLCSHL 500
```

```
RESULT 41
US-10-104-047-3399
```

```

; Sequence 3399, Application US/10104047
; Patent No. 6943241
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 6943241el full length cDNA
; FILE REFERENCE: HI-A0105
; CURRENT APPLICATION NUMBER: US/10/104,047
; PRIOR FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3399
; LENGTH: 728
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-104-047-3399

```

```

Query Match      70.8%; Score 34; DB 2; Length 728;
Best Local Similarity 75.0%; Pred. No. 3.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      2 LKPOLCTEL 9
DB      493 LKPOLCSHL 500

```

```

RESULT 42
US-09-252-991A-31445
; Sequence 31445, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 31445
; LENGTH: 163
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-31445

```

```

Query Match      68.8%; Score 33; DB 2; Length 163;
Best Local Similarity 71.4%; Pred. No. 1.2e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

```

```

QY      1 KLPOLCT 7
DB      36 RLPELCT 42

```

```

RESULT 43
US-09-248-796A-16504
; Sequence 16504, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13

```

```

; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 16504
; LENGTH: 304
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-16504

```

```

Query Match      68.8%; Score 33; DB 2; Length 304;
Best Local Similarity 85.7%; Pred. No. 2.3e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

```

```

QY      1 KLPOLCTEL 9
DB      172 KLPOLCRDI 180

```

```

RESULT 44
US-09-369-247-60
; Sequence 60, Application US/09369247
; Patent No. 6569992
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 44 Human Secreted Proteins
; FILE REFERENCE: P2024P1
; CURRENT APPLICATION NUMBER: US/09/369,247
; CURRENT FILING DATE: 1999-08-05
; EARLIER APPLICATION NUMBER: 60/074,118
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,157
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,137
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,341
; EARLIER FILING DATE: 1998-02-09
; EARLIER APPLICATION NUMBER: 60/074,141
; EARLIER FILING DATE: 1998-02-09
; NUMBER OF SEQ ID NOS: 172
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 60
; LENGTH: 308
; TYPE: PRT
; ORGANISM: Homo sapiens

```

```

; FEATURE:
; NAME/KEY: SITE
; LOCATION: (165)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (247)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (308)
; OTHER INFORMATION: Xaa equals stop translation
US-09-369-247-60

```

```

Query Match      68.8%; Score 33; DB 2; Length 308;
Best Local Similarity 85.7%; Pred. No. 2.3e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      3 POLCTEL 9
DB      286 PPLCTEL 292

```

```

RESULT 45
US-10-062-548-60
; Sequence 60, Application US/10062548
; Patent No. 6924356
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: 44 Human Secreted Proteins
; FILE REFERENCE: P2024P1

```

```

; CURRENT APPLICATION NUMBER: US/10/062,548
; CURRENT FILING DATE: 2002-02-05
; PRIOR APPLICATION NUMBER: 09/369,247
; PRIOR FILING DATE: 1999-08-05
; PRIOR APPLICATION NUMBER: 60/074,118
; PRIOR FILING DATE: 1998-02-09
; PRIOR APPLICATION NUMBER: 60/074,157
; PRIOR FILING DATE: 1998-02-09
; PRIOR APPLICATION NUMBER: 60/074,137
; PRIOR FILING DATE: 1998-02-09
; PRIOR APPLICATION NUMBER: 60/074,341
; PRIOR FILING DATE: 1998-02-09
; PRIOR APPLICATION NUMBER: 60/074,141
; PRIOR FILING DATE: 1998-02-09
; NUMBER OF SEQ ID NOS: 172
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO: 60
; LENGTH: 308
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (165)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (247)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (308)
; OTHER INFORMATION: Xaa equals stop translation
; US-10-062-548-60

Query Match      68.8%; Score 33; DB 2; Length 308;
Best Local Similarity 85.7%; Pred. No. 2.3e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      3 POLCTEL 9
Db      286 PDLCTEL 292

RESULT 46
US-10-104-047-3188
; Sequence 3188, Application US/10104047
; Patent No. 6943241
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: NO. 6943241el full length cDNA
; FILE REFERENCE: H1-A0105
; CURRENT APPLICATION NUMBER: US/10/104,047
; CURRENT FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; PRIOR FILING DATE:
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO: 3188
; LENGTH: 384
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-10-104-047-3188

Query Match      68.8%; Score 33; DB 2; Length 384;
Best Local Similarity 75.0%; Pred. No. 2.9e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy      2 LPOLCTEL 9
Db      348 LPSLCTSL 355

RESULT 47
US-09-543-681A-7117
; Sequence 7117, Application US/09543681A
```

```

; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETTON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; CURRENT FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO: 7117
; LENGTH: 476
; TYPE: PRT
; ORGANISM: Proteus mirabilis
; US-09-543-681A-7117

Query Match      68.8%; Score 33; DB 2; Length 476;
Best Local Similarity 85.7%; Pred. No. 3.5e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy      3 POLCTEL 9
Db      46 PDLCTEL 52

RESULT 48
US-08-551-211-3
; Sequence 3, Application US/08551211
; Patent No. 5843735
; GENERAL INFORMATION:
; APPLICANT: LEE, Jung Joon
; APPLICANT: KIM, Young Ho
; APPLICANT: HONG, Soon Kwang
; APPLICANT: HONG, Young Soo
; APPLICANT: HWANG, Cheol Kyu
; APPLICANT: KIM, Hang Sub
; TITLE OF INVENTION: AKAVINONE C-11 HYDROXYLASE, GENE CODING
; TITLE OF INVENTION: FOR SAME, EXPRESSION VECTOR THEREFOR, AND
; TITLE OF INVENTION: PROCESS FOR PREPARING HYBRID ANTIBIOTICS
; TITLE OF INVENTION: BY USING SAID VECTOR
; NUMBER OF SEQUENCES: 5
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: LEE, Jung Joon
; STREET: Hanbit Apt. 132-201, Oeun-dong, Youseong-ku
; CITY: Daejeon
; STATE: Daejeon
; COUNTRY: Republic of Korea
; ZIP: 305-333
; ADDRESSEE: KIM, Young Ho
; STREET: Hanbit Apt. 125-1504, Oeun-dong, Youseong-ku
; CITY: Daejeon
; STATE: Daejeon
; COUNTRY: Republic of Korea
; ZIP: 305-333
; ADDRESSEE: HONG, Soon Kwang
; STREET: #231-32 Mochung-dong
; CITY: Cheongju-si
; STATE: Chungcheongbuk-do
; COUNTRY: Republic of Korea
; ZIP: 360-140
; ADDRESSEE: HONG, Young Soo
; STREET: #San-1, Oeun-dong, Youseong-ku
; CITY: Daejeon
; STATE: Daejeon
; COUNTRY: Republic of Korea
; ZIP: 305-333
; ADDRESSEE: HWANG, Cheol Kyu
; STREET: #San-1, Oeun-dong, Youseong-ku
; CITY: Daejeon
; STATE: Daejeon
; COUNTRY: Republic of Korea
; ZIP: 305-333
```

```

; ADDRESSEE: KIM, Hang Sub
; STREET: Sindonga Apt. 11-1403, Yongjeon-dong, Dong-ku
; CITY: Daejeon
; STATE: Daejeon
; COUNTRY: Republic of Korea
; ZIP: 300-200
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.5 inch, 1.44 MB storage
; COMPUTER: IBM PC/AT
; OPERATING SYSTEM: MS-DOS
; SOFTWARE: Word Perfect 5.1
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/551,211
; FILING DATE: 31-OCT-1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: KR 95-1950
; FILING DATE: 2-MAR-1995
; ATTORNEY/AGENT INFORMATION:
; NAME: Shahar Islam
; REGISTRATION NUMBER: 32,507
; REFERENCE/DOCKET NUMBER: DT-1421
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212) 278-1000
; TELEFAX: (212) 953-7249
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 536 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-551-211-3

Query Match      68.8%; Score 33; DB 1; Length 536;
Best Local Similarity 85.7%; Pred. No. 4e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      3 POLCTEL 9
Db      262 PORCTEL 268

```

```

RESULT 49
US-09-673-198-1
; Sequence 1, Application US/09673198
; Patent No. 6806076
; GENERAL INFORMATION:
; APPLICANT: MIYAKE, Koichiro, HASHIMOTO, Shinichi, MOTOMYAMA Hiroaki,
; APPLICANT: OZAKI, Akiyo, SETO, Haruo, KUZAYAMA, Tomohisa, TAKAHASHI, Shunji
; TITLE OF INVENTION: A process for producing isoprenoid compounds by
; TITLE OF INVENTION: microorganisms and a method for screening compounds with
; TITLE OF INVENTION: antibiotic or weeding activity
; FILE REFERENCE:
; CURRENT APPLICATION NUMBER: US/09/673,198
; CURRENT FILING DATE: 2000-10-12
; PRIOR APPLICATION NUMBER: JP98/103101
; PRIOR FILING DATE: 1998-04-14
; PRIOR APPLICATION NUMBER: JP98/221910
; PRIOR FILING DATE: 1998-08-05
; PRIOR APPLICATION NUMBER: JP99/035739
; PRIOR FILING DATE: 1999-02-15
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: Patent Ver. 2.0
; SEQ ID NO 1
; LENGTH: 620
; TYPE: PRT
; ORGANISM: Escherichia coli
; US-09-673-198-1

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Query Match      68.8%; Score 33; DB 2; Length 620;
Best Local Similarity 75.0%; Pred. No. 4.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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QY      2 LPOLCTEL 9
Db      28 LPRCTDEL 35

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; Sequence 13113, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 13113
; LENGTH: 626
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
; US-09-489-039A-13113

Query Match      68.8%; Score 33; DB 2; Length 626;
Best Local Similarity 75.0%; Pred. No. 4.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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Db      34 LPRCTDEL 41

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OM protein - protein search, using sw model

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Perfect score: 48
Sequence: 1 KLPOLCTEL 9

Scoring table: BLOSUM62
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Searched: 1867569 seqs, 417829326 residues

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Listing first 1000 summaries

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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8	48	100.0	9	5	US-10-751-845-55 Sequence 55, Appli
9	48	100.0	12	6	US-11-021-949-77 Sequence 7, Appli
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11	48	100.0	20	5	US-10-751-845-64 Sequence 64, Appli
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452	31	64.6	421	5	US-10-739-930-10809	Sequence 10809, A	525	31	64.6	1214	4	US-10-437-963-129932	Sequence 129932, A
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454	31	64.6	437	5	US-10-739-930-3381	Sequence 7381, Appl1	527	31	64.6	3084	4	US-10-262-670-2	Sequence 2, Appl1
455	31	64.6	442	3	US-09-815-242-12307	Sequence 12307, A	528	30	62.5	10	4	US-10-182-252A-330	Sequence 390, App
456	31	64.6	452	3	US-09-996-620-16	Sequence 16, Appl1	529	30	62.5	10	4	US-10-444-534-1	Sequence 1, Appl1
457	31	64.6	462	6	US-11-066-236-16	Sequence 16, Appl1	530	30	62.5	30	3	US-10-424-599-24926	Sequence 240926, A
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555	30	62.5	83	4	US-10-127-033-112	Sequence 112, App	628	30	62.5	256	3	US-09-815-442-11725	Sequence 11725, App
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572	30	62.5	111	4	US-10-437-963-201627	Sequence 201627, App	645	30	62.5	326	4	US-10-437-963-122213	Sequence 29, App1
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576	30	62.5	119	4	US-10-437-963-156051	Sequence 156051, App	649	30	62.5	328	4	US-10-424-599-473537	Sequence 39990, App
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582	30	62.5	129	4	US-10-424-599-261473	Sequence 261473, App	655	30	62.5	363	5	US-10-369-493-15938	Sequence 15938, App
583	30	62.5	130	5	US-10-511-698-12	Sequence 12, App1	656	30	62.5	367	4	US-10-762-769-8	Sequence 7, App1
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591	30	62.5	157	4	US-10-425-115-345034	Sequence 345034, App	664	30	62.5	388	6	US-11-108-597-9	Sequence 9, App1
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594	30	62.5	161	5	US-10-511-698-11	Sequence 11, App1	667	30	62.5	391	4	US-10-210-281-48	Sequence 48, App1
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596	30	62.5	176	5	US-10-764-833-29	Sequence 29, App1	669	30	62.5	396	4	US-10-424-599-248725	Sequence 248725, App
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598	30	62.5	186	4	US-10-425-114-44149	Sequence 44149, App	671	30	62.5	421	4	US-10-424-599-248725	Sequence 248725, App
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610	30	62.5	224	5	US-10-872-155-397	Sequence 397, App	683	30	62.5	463	4	US-10-732-923-2055	Sequence 2055, App
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688	30	62.5	498	5	US-10-866-484-2	Sequence 2, Appl1	761	30	62.5	877	4	US-10-243-023-90	Sequence 90, Appl
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704	30	62.5	567	4	US-10-334-322A-62	Sequence 62, Appl	777	30	62.5	877	4	US-10-238-325-90	Sequence 90, Appl
705	30	62.5	567	4	US-10-646-640-11	Sequence 11, Appl	778	30	62.5	877	4	US-10-238-326-90	Sequence 90, Appl
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712	30	62.5	567	5	US-10-511-698-10	Sequence 10, Appl	785	30	62.5	877	4	US-10-243-388-90	Sequence 90, Appl
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716	30	62.5	592	3	US-09-917-788-5	Sequence 5, Appl1	789	30	62.5	877	4	US-10-245-079-90	Sequence 90, Appl
717	30	62.5	604	3	US-09-738-626-3886	Sequence 3886, Ap	790	30	62.5	877	4	US-10-245-127-90	Sequence 90, Appl
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723	30	62.5	722	4	US-10-444-467-8	Sequence 8, Appl1	796	30	62.5	877	4	US-10-245-878-90	Sequence 90, Appl
724	30	62.5	722	4	US-10-424-599-161076	Sequence 161076, Appl	797	30	62.5	877	4	US-10-245-899-90	Sequence 90, Appl
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729	30	62.5	739	4	US-10-717-282-12	Sequence 12, Appl	802	30	62.5	877	4	US-10-237-471-90	Sequence 90, Appl
730	30	62.5	741	4	US-10-320-797-3278	Sequence 3278, Ap	803	30	62.5	877	4	US-10-238-261-90	Sequence 90, Appl
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732	30	62.5	798	6	US-11-097-143-11032	Sequence 11032, A	805	30	62.5	877	4	US-10-241-860-90	Sequence 90, Appl
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ALIGNMENTS

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; Sequence 101, Application US/09909460
; Publication No. US20020182258A1
; GENERAL INFORMATION:
; APPLICANT: Lunsford, Lynn B.
; APPLICANT: Putnam, David
; APPLICANT: Hedley, Mary Lynn
; TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY OF NUCLEIC
; TITLE OF INVENTION: ACID
; FILE REFERENCE: 08191/014001
; CURRENT APPLICATION NUMBER: US/09/909,460
; CURRENT FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/321,346
; PRIOR FILING DATE: EARLIER FILING DATE: 1999-05-27
; NUMBER OF SEQ ID NOS: 114
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 101
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Hepatitis B virus
US-09-909-460-101

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Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 2
US-09-872-836-101
; Sequence 101, Application US/09872836
; Publication No. US20040142475A1
; GENERAL INFORMATION:
; APPLICANT: Bartman, Shikha P.
; APPLICANT: McKeever, Una
; APPLICANT: Hedley, Mary Lynne
; TITLE OF INVENTION: DELIVERY SYSTEMS FOR BIOACTIVE AGENTS
; FILE REFERENCE: 08191-018001
; CURRENT APPLICATION NUMBER: US/09/872,836
; CURRENT FILING DATE: 2001-06-01
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; PRIOR APPLICATION NUMBER: US 60/208,830
; PRIOR FILING DATE: 2000-06-02
; NUMBER OF SEQ ID NOS: 120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 101
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-872-836-101

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; GENERAL INFORMATION:
; APPLICANT: Delisi, Charles
; APPLICANT: Bezotsky, Jay
; APPLICANT: Gulikota, Kamalakara
; APPLICANT: Vaccaro, Dennis
; APPLICANT: Weng, Zhiping
; APPLICANT: Zhang, Chao
; TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
; TITLE OF INVENTION: COMPOSITIONS THEREOF
; FILE REFERENCE: BU-035AX
; CURRENT APPLICATION NUMBER: US/10/133,210
; CURRENT FILING DATE: 2002-04-26
; NUMBER OF SEQ ID NOS: 281
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; Publication No. US20040132088A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Diamond, David C.
; APPLICANT: Qiu, Zhiyong
; APPLICANT: Lei, Xiang-Dong
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
; TITLE OF INVENTION: TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
; FILE REFERENCE: MANKR.022C1
; CURRENT APPLICATION NUMBER: US/10/777,053
; CURRENT FILING DATE: 2004-02-10
; PRIOR APPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: 60/336,968
; PRIOR FILING DATE: 2001-11-07
; NUMBER OF SEQ ID NOS: 979
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US-10-777-053-546
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; Sequence 546, Application US/10837217
; Publication No. US20040203051A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Diamond, David C.
; APPLICANT: Qiu, Xiang-Dong
; APPLICANT: Lei, Xiang-Dong
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
; TITLE OF INVENTION: TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
; FILE REFERENCE: MANMK 022C2
; CURRENT APPLICATION NUMBER: US/10/837,217
; CURRENT FILING DATE: 2004-04-30
; PRIOR APPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: 60/336,968
; PRIOR FILING DATE: 2001-11-07
; NUMBER OF SEQ ID NOS: 979
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 546
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-10-837-217-546
```

```
Query Match          100.0%; Score 48; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
    |||||
Db 1 KLPOLCTEL 9
```

```
RESULT 6
US-10-758-970-101
; Sequence 101, Application US/10758970
; Publication No. US20050037086A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Hsu, Yung-Yueh
; APPLICANT: Tyo, Michael
; TITLE OF INVENTION: CONTINUOUS-FLOW METHOD FOR PREPARING MICROPARTICLES
; FILE REFERENCE: 08191-012001
; CURRENT APPLICATION NUMBER: US/10/758,970
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: US/09/715,708A
; PRIOR FILING DATE: 2000-11-17
; PRIOR APPLICATION NUMBER: US 60/166,516
; PRIOR FILING DATE: 1999-11-19
; NUMBER OF SEQ ID NOS: 109
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 101
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human papilloma virus
```

```
US-10-758-970-101
```

```
Query Match          100.0%; Score 48; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
    |||||
Db 1 KLPOLCTEL 9
```

```
RESULT 7
US-10-484-063-1
; Sequence 1, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILTERMO
; APPLICANT: POLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-1
```

```
Query Match          100.0%; Score 48; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
    |||||
Db 1 KLPOLCTEL 9
```

```
RESULT 8
US-10-751-845-55
; Sequence 55, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Chiciz, Roman M.
; APPLICANT: Urban, Robert G.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 55
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-55
```

```
Query Match          100.0%; Score 48; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
```


Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
| | | | |
Db 1 KLPOLCTEL 9

RESULT 9
US-11-021-949-7
; Sequence 7, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 7
; LENGTH: 12
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-7

Query Match 100.0%; Score 48; DB 6; Length 12;
Best Local Similarity 100.0%; Pred. No. 0.072;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
| | | | |
Db 4 KLPOLCTEL 12

RESULT 10
US-10-476-570-20
; Sequence 20, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: PR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 20
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 17-31
US-10-476-570-20

Query Match 100.0%; Score 48; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.089;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
| | | | |
Db 2 KLPOLCTEL 10

RESULT 11
US-10-751-845-64
; Sequence 64, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chiciz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 64
; LENGTH: 20
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-64

Query Match 100.0%; Score 48; DB 5; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
| | | | |
Db 12 KLPOLCTEL 20

RESULT 12
US-10-476-570-8
; Sequence 8, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: PR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 8
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-34
US-10-476-570-8

Query Match 100.0%; Score 48; DB 4; Length 21;

Best Local Similarity 100.0%; Pred. No. 0.12; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 KLPOCTEL 9
Db 5 KLPOCTEL 13

RESULT 13
US-11-021-949-1

/ Sequence 1, Application US/11021949
/ Publication No. US20050142541A1
/ GENERAL INFORMATION:
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ TITLE OF INVENTION: AND METHODS OF THEIR USE
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
/ CURRENT FILING DATE: 2004-12-23
/ PRIOR APPLICATION NUMBER: 60/532,373
/ PRIOR FILING DATE: 2003-12-23
/ NUMBER OF SEQ ID NOS: 361
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 1
/ LENGTH: 25
/ TYPE: PRT
/ ORGANISM: human papilloma virus (HPV)
US-11-021-949-1

Query Match 100.0%; Score 48; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.15; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 KLPOCTEL 9
Db 10 KLPOCTEL 18

RESULT 14
US-10-476-570-53

/ Sequence 53, Application US/10476570
/ Publication No. US20040170644A1
/ GENERAL INFORMATION:
/ APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
/ APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
/ APPLICANT: MAILLIERE, Bernard
/ APPLICANT: BOURGAULT-VILLADA, Isabelle
/ APPLICANT: GUILLET, Jean-Gerard
/ TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
/ TITLE OF INVENTION: Papillomavirus proteins and uses thereof
/ FILE REFERENCE: 45636-5071-US
/ CURRENT APPLICATION NUMBER: US/10/476,570
/ CURRENT FILING DATE: 2003-11-04
/ PRIOR APPLICATION NUMBER: PCT/FR02/01533
/ PRIOR FILING DATE: 2002-05-03
/ PRIOR APPLICATION NUMBER: FR 01 05980
/ PRIOR FILING DATE: 2001-05-04
/ NUMBER OF SEQ ID NOS: 63
/ SOFTWARE: PatentIn Ver. 2.1
/ SEQ ID NO 53
/ LENGTH: 30
/ TYPE: PRT
/ ORGANISM: artificial sequence
/ FEATURE:
/ OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53

Query Match 100.0%; Score 48; DB 4; Length 30;

Best Local Similarity 100.0%; Pred. No. 0.17; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 KLPOCTEL 9
Db 4 KLPOCTEL 12

RESULT 15
US-10-858-384-4

/ Sequence 4, Application US/10858384
/ Publication No. US20050033025A1
/ GENERAL INFORMATION:
/ APPLICANT: CHOPPIN, JEANNINE
/ APPLICANT: BOURGAULT VILLADA, ISABELLE
/ APPLICANT: GUILLET, JEAN-GERARD
/ APPLICANT: CONNAN, FRANCES
/ APPLICANT: FERRIES, ESTELLE
/ TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
/ TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
/ FILE REFERENCE: 0508-1037-1
/ CURRENT APPLICATION NUMBER: US/10/858,384
/ CURRENT FILING DATE: 2004-06-02
/ PRIOR APPLICATION NUMBER: FR 9907012
/ PRIOR FILING DATE: 1999-06-03
/ NUMBER OF SEQ ID NOS: 24
/ SOFTWARE: PatentIn Ver. 3.2
/ SEQ ID NO 4
/ LENGTH: 30
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match 100.0%; Score 48; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.17; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 KLPOCTEL 9
Db 4 KLPOCTEL 12

RESULT 16
US-10-476-570-9

/ Sequence 9, Application US/10476570
/ Publication No. US20040170644A1
/ GENERAL INFORMATION:
/ APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
/ APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
/ APPLICANT: MAILLIERE, Bernard
/ APPLICANT: BOURGAULT-VILLADA, Isabelle
/ APPLICANT: GUILLET, Jean-Gerard
/ TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
/ TITLE OF INVENTION: Papillomavirus proteins and uses thereof
/ FILE REFERENCE: 45636-5071-US
/ CURRENT APPLICATION NUMBER: US/10/476,570
/ CURRENT FILING DATE: 2003-11-04
/ PRIOR APPLICATION NUMBER: PCT/FR02/01533
/ PRIOR FILING DATE: 2002-05-03
/ PRIOR APPLICATION NUMBER: FR 01 05980
/ PRIOR FILING DATE: 2001-05-04
/ NUMBER OF SEQ ID NOS: 63
/ SOFTWARE: PatentIn Ver. 2.1
/ SEQ ID NO 9
/ LENGTH: 32
/ TYPE: PRT
/ ORGANISM: artificial sequence
/ FEATURE:
/ OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-9

Query Match 100.0%; Score 48; DB 4; Length 30;

```
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9
Query Match          100.0%; Score 48; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.19;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 5 KLPOLCTEL 13

RESULT 17
US-10-476-570-19
; Sequence 19, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUEVILLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OR INVENTION: Papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 33
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19
Query Match          100.0%; Score 48; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.19;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 5 KLPOLCTEL 13

RESULT 18
US-10-751-845-126
; Sequence 126, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 126
; LENGTH: 117

; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-126
Query Match          100.0%; Score 48; DB 5; Length 117;
Best Local Similarity 100.0%; Pred. No. 0.64;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 12 KLPOLCTEL 20

RESULT 19
US-10-177-390-6
; Sequence 6, Application US/10177390
; Publication No. US20030143743A1
; GENERAL INFORMATION:
; APPLICANT: Schuler, Gerold
; APPLICANT: N.V. Antwerp Innovatiecentrum
; TITLE OF INVENTION: Improved transfection of Eucaryotic Cells with linear
; TITLE OF INVENTION: Polynucleotides by Electroporation
; FILE REFERENCE: 021505w/JH/ml
; CURRENT APPLICATION NUMBER: US/10/177,390
; CURRENT FILING DATE: 2002-06-20
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-177-390-6
Query Match          100.0%; Score 48; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.82;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 11 KLPOLCTEL 19

RESULT 20
US-10-484-063-20
; Sequence 20, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILERMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-20
Query Match          100.0%; Score 48; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.82;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 KLPOLCTEL 9
| | | | |
Db 11 KLPOLCTEL 19

RESULT 21
US-10-484-063-27
; Sequence 27, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILLEMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
; FILE REFERENCE: UISC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 27
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match 100.0%; Score 48; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.82;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
| | | | |
Db 11 KLPOLCTEL 19

RESULT 22
US-10-858-384-2
; Sequence 2, Application US/10858384
; Publication No. US20050033025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOUGAUDIT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match 100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.86;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
| | | | |
Db 18 KLPOLCTEL 26

RESULT 23
US-10-367-057-16
; Sequence 16, Application US/10367057
; Publication No. US20050100554A1
; GENERAL INFORMATION:
; APPLICANT: Cuthill, Scott;
; APPLICANT: Jackson, Amanda;
; APPLICANT: Lewin, David A.;
; APPLICANT: Ooi, Chean Eng
; TITLE OF INVENTION: Complexes and Methods of Using Same
; FILE REFERENCE: 21402-559
; CURRENT APPLICATION NUMBER: US/10/367,057
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: 60/256,911
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 198
; SOFTWARE: CirusSeqList version 0.1
; SEQ ID NO 16
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-367-057-16

Query Match 100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.86;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
| | | | |
Db 18 KLPOLCTEL 26

RESULT 24
US-11-021-949-13
; Sequence 13, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SABIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match 100.0%; Score 48; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.86;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
| | | | |
Db 18 KLPOLCTEL 26

RESULT 25
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel

```
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; CURRENT FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03771
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Construct
US-10-472-724-2
```

```
Query Match          100.0%; Score 48; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 0.93;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 23 KLPOLCTEL 31
```

```
RESULT 26
US-10-751-845-157
; Sequence 157, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 157
; LENGTH: 236
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-157
```

```
Query Match          100.0%; Score 48; DB 5; Length 236;
Best Local Similarity 100.0%; Pred. No. 1.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 12 KLPOLCTEL 20
```

```
RESULT 27
US-10-751-845-158
; Sequence 158, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
```

```
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 158
; LENGTH: 237
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-158
```

```
Query Match          100.0%; Score 48; DB 5; Length 237;
Best Local Similarity 100.0%; Pred. No. 1.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 13 KLPOLCTEL 21
```

```
RESULT 28
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOUL, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTI-TUMORAL COMPOSITION BASED ON IMMUNOGENIC
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; CURRENT FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-11-072-288-1
```

```
Query Match          100.0%; Score 48; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 1.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPOLCTEL 9
Db 46 KLPOLCTEL 54
```

```
RESULT 29
US-10-751-845-160
; Sequence 160, Application US/10751845
; Publication No. US20050100928A1
```

```

; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 160
; LENGTH: 261
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-160

Query Match          100.0%; Score 48; DB 5; Length 261;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KLPOLCTEL 9
Db 37 KLPOLCTEL 45

RESULT 30
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Publication No. US20020081329A1
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; CURRENT FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match          100.0%; Score 48; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KLPOLCTEL 9
Db 18 KLPOLCTEL 26

RESULT 31
US-10-000-903-4
; Sequence 4, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
```

```

; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP96/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-4

Query Match          100.0%; Score 48; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KLPOLCTEL 9
Db 124 KLPOLCTEL 132

RESULT 32
US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimaeic protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4

Query Match          100.0%; Score 48; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 KLPOLCTEL 9
Db 124 KLPOLCTEL 132

RESULT 33
US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
```

```
/ APPLICANT: Delisse, Anne-Marie Eva Fernande
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ APPLICANT: Lombardo-Bencheikh, Angela
/ TITLE OF INVENTION: Vaccine
/ FILE REFERENCE: B45107
/ CURRENT APPLICATION NUMBER: US/10/000,903
/ CURRENT FILING DATE: 2001-10-01
/ PRIOR APPLICATION NUMBER: PCT/EP98/05285
/ PRIOR FILING DATE: 1998-08-17
/ PRIOR APPLICATION NUMBER: GB 9717953.5
/ PRIOR FILING DATE: 1997-08-22
/ NUMBER OF SEQ ID NOS: 23
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 10
/ LENGTH: 292
/ TYPE: PRT
/ ORGANISM: Homo sapien
US-10-000-903-10
```

```
Query Match          100.0%; Score 48; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
    |||||
Db 143 KLPOLCTEL 151
```

RESULT 34

```
US-10-899-771-10
/ Sequence 10, Application US/10899771
/ Publication No. US20050031638A1
/ GENERAL INFORMATION:
/ APPLICANT: Dalemans, Wilfried L.J.
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
/ TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
/ FILE REFERENCE: B45124
/ CURRENT APPLICATION NUMBER: US/10/899,771
/ CURRENT FILING DATE: 2004-07-27
/ PRIOR APPLICATION NUMBER: US/09/581,976
/ PRIOR FILING DATE: 2000-06-20
/ PRIOR APPLICATION NUMBER: PCT/EP98/08563
/ PRIOR FILING DATE: 1998-12-18
/ PRIOR APPLICATION NUMBER: GB 9727262.9
/ PRIOR FILING DATE: 1997-12-24
/ NUMBER OF SEQ ID NOS: 28
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 10
/ LENGTH: 292
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Chimaeric protein (Clyta from Streptococcus
/ OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type
/ OTHER INFORMATION: 16)
US-10-899-771-10
```

```
Query Match          100.0%; Score 48; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
    |||||
Db 143 KLPOLCTEL 151
```

```
RESULT 35
US-10-000-903-6
/ Sequence 6, Application US/10000903
/ Publication No. US20020182221A1
/ GENERAL INFORMATION:
/ APPLICANT: Bruck, Claudine
```

```
/ APPLICANT: Cabezon Silva, Teresa
/ APPLICANT: Delisse, Anne-Marie Eva Fernande
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ APPLICANT: Lombardo-Bencheikh, Angela
/ TITLE OF INVENTION: Vaccine
/ FILE REFERENCE: B45107
/ CURRENT APPLICATION NUMBER: US/10/000,903
/ CURRENT FILING DATE: 2001-10-01
/ PRIOR APPLICATION NUMBER: PCT/EP98/05285
/ PRIOR FILING DATE: 1998-08-17
/ PRIOR APPLICATION NUMBER: GB 9717953.5
/ PRIOR FILING DATE: 1997-08-22
/ NUMBER OF SEQ ID NOS: 23
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 6
/ LENGTH: 371
/ TYPE: PRT
/ ORGANISM: Homo sapien
US-10-000-903-6
```

```
Query Match          100.0%; Score 48; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
    |||||
Db 124 KLPOLCTEL 132
```

RESULT 36

```
US-10-899-771-6
/ Sequence 6, Application US/10899771
/ Publication No. US20050031638A1
/ GENERAL INFORMATION:
/ APPLICANT: Dalemans, Wilfried L.J.
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
/ TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
/ FILE REFERENCE: B45124
/ CURRENT APPLICATION NUMBER: US/10/899,771
/ CURRENT FILING DATE: 2004-07-27
/ PRIOR APPLICATION NUMBER: US/09/581,976
/ PRIOR FILING DATE: 2000-06-20
/ PRIOR APPLICATION NUMBER: PCT/EP98/08563
/ PRIOR FILING DATE: 1998-12-18
/ PRIOR APPLICATION NUMBER: GB 9727262.9
/ PRIOR FILING DATE: 1997-12-24
/ NUMBER OF SEQ ID NOS: 28
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 6
/ LENGTH: 371
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
/ OTHER INFORMATION: influenzae B and E6E7 fusion from Human papilloma
/ OTHER INFORMATION: virus type 16)
US-10-899-771-6
```

```
Query Match          100.0%; Score 48; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 KLPOLCTEL 9
    |||||
Db 124 KLPOLCTEL 132
```

```
RESULT 37
US-10-000-903-14
/ Sequence 14, Application US/10000903
/ Publication No. US20020182221A1
/ GENERAL INFORMATION:
```

```

; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benckelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-14

Query Match          100.0%; Score 48; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 143 KLPOLCTEL 151

RESULT 38
US-10-899-771-14
; Sequence 14, Application US/10899771
; Publication No. US20050031636A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuncted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimaeic protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6B7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-14

Query Match          100.0%; Score 48; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 143 KLPOLCTEL 151

RESULT 39
US-10-367-095-10
; Sequence 10, Application US/10367095
; Publication No. US20030228696A1
```

```

; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
; FILE REFERENCE: 44149-10S1
; CURRENT APPLICATION NUMBER: US/10/367,095
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-095-10

Query Match          100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 488 KLPOLCTEL 496

RESULT 40
US-10-368-046-10
; Sequence 10, Application US/10368046
; Publication No. US20040063188A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; APPLICANT: Victoria Cioce
; TITLE OF INVENTION: Method for Isolation and Purification of
; TITLE OF INVENTION: Expressed Gene Products In Vitro
; FILE REFERENCE: 44149-30S1
; CURRENT APPLICATION NUMBER: US/10/368,046
; CURRENT FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
```

;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,135
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-368-046-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 488 KLPOLCTEL 496

RESULT 41
US-10-367-367-10
;; Sequence 10, Application US/10367367
;; Publication No. US20040121465A1
;; GENERAL INFORMATION:
;; APPLICANT: Robin A. Robinson
;; TITLE OF INVENTION: Optimization of Gene Sequences of
;; TITLE OF INVENTION: Virus-Like Particles for Expression in Insect Cells
;; FILE REFERENCE: 44149-2US1
;; CURRENT APPLICATION NUMBER: US/10/367,367
;; CURRENT FILING DATE: 2003-02-15
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-367-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 488 KLPOLCTEL 496

RESULT 42
US-10-918-337-10
;; Sequence 10, Application US/10918337
;; Publication No. US20050118191A1
;; GENERAL INFORMATION:
;; APPLICANT: NOVAVAX, INC., et al.

;; TITLE OF INVENTION: Optimization of Gene Sequences of
;; TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
;; FILE REFERENCE: 19065/2132
;; CURRENT APPLICATION NUMBER: US/10/918,337
;; CURRENT FILING DATE: 2004-08-13
;; PRIOR APPLICATION NUMBER: PCT/US03/04473
;; PRIOR FILING DATE: 2003-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10

Query Match 100.0%; Score 48; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 488 KLPOLCTEL 496

RESULT 43
US-10-472-661-1
;; Sequence 1, Application US/10472661
;; Publication No. US20040106551A1
;; GENERAL INFORMATION:
;; APPLICANT: Knleif, Samir N.
;; APPLICANT: Berzofsky, Jay A.
;; TITLE OF INVENTION: HUMAN PAPILLOMA VIRUS IMMUNOREACTIVE
;; TITLE OF INVENTION: PEPTIDES
;; FILE REFERENCE: 14014, 0406U2
;; CURRENT APPLICATION NUMBER: US/10/472,661
;; CURRENT FILING DATE: 2003-09-22
;; PRIOR APPLICATION NUMBER: PCT/US02/09261
;; PRIOR FILING DATE: 2002-03-22
;; PRIOR APPLICATION NUMBER: 60/278,520
;; PRIOR FILING DATE: 2001-03-23
;; NUMBER OF SEQ ID NOS: 9
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 1
;; LENGTH: 9
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: Description of Artificial Sequence; note =
US-10-472-661-1

Query Match 89.6%; Score 43; DB 4; Length 9;

Best Local Similarity 88.9%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 1 KLPDLCTEL 9

RESULT 44

US-10-751-845-124
; Sequence 124, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chiciz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; PRIOR FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 124
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-124

Query Match 89.6%; Score 43; DB 5; Length 9;
Best Local Similarity 88.9%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 1 KLPDLCTEL 9

RESULT 45

US-11-021-949-8
; Sequence 8, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 12
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-8

Query Match 89.6%; Score 43; DB 6; Length 12;
Best Local Similarity 88.9%; Pred. No. 0.58;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9

Db 4 KLPDLCTEL 12

RESULT 46
US-11-021-949-60
; Sequence 60, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 60
; LENGTH: 12
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-60

Query Match 89.6%; Score 43; DB 6; Length 12;
Best Local Similarity 88.9%; Pred. No. 0.58;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 4 KLPDLCTEL 12

RESULT 47
US-11-021-949-2
; Sequence 2, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-2

Query Match 89.6%; Score 43; DB 6; Length 25;
Best Local Similarity 88.9%; Pred. No. 1.2;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOLCTEL 9
Db 10 KLPDLCTEL 18

RESULT 48

```
US-11-021-949-57
; Sequence 57, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 57
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-57

Query Match      89.6%; Score 43; DB 6; Length 25;
Best Local Similarity 88.9%; Pred. No. 1.2;
Matches      8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 KLPOLCTEL 9
Db      10 KLPDLCTEL 18

RESULT 49
US-10-751-845-152
; Sequence 152, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Robert M.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 152
; LENGTH: 42
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-152

Query Match      89.6%; Score 43; DB 5; Length 42;
Best Local Similarity 88.9%; Pred. No. 1.9;
Matches      8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 KLPOLCTEL 9
Db      5 KLPDLCTEL 13

RESULT 50
US-10-751-845-159
; Sequence 159, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
```

```
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 159
; LENGTH: 119
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-159

Query Match      89.6%; Score 43; DB 5; Length 119;
Best Local Similarity 88.9%; Pred. No. 5.3;
Matches      8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 KLPOLCTEL 9
Db      5 KLPDLCTEL 13
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Search completed: May 5, 2006, 08:50:29
Job time : 62.3 secs

GenCore version 5.1.7
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OM protein - protein search, using SW model

Run on: May 5, 2006, 08:40:52 / Search time 8.4 Seconds
(without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-2
Perfect score: 48
Sequence: 1 KLPOLCTEL 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :
1: /SIDS5/ptcodata/1/pubppaa/US08_NEW_PUB.pep1.*
2: /SIDS5/ptcodata/1/pubppaa/US06_NEW_PUB.pep.*
3: /SIDS5/ptcodata/1/pubppaa/US07_NEW_PUB.pep.*
4: /SIDS5/ptcodata/1/pubppaa/US08_NEW_PUB.pep.*
5: /SIDS5/ptcodata/1/pubppaa/PCT_NEW_PUB.pep.*
6: /SIDS5/ptcodata/1/pubppaa/US09_NEW_PUB.pep.*
7: /SIDS5/ptcodata/1/pubppaa/US09_NEW_PUB.pep1.*
8: /SIDS5/ptcodata/1/pubppaa/US10_NEW_PUB.pep.*
9: /SIDS5/ptcodata/1/pubppaa/US10_NEW_PUB.pep1.*
10: /SIDS5/ptcodata/1/pubppaa/US11_NEW_PUB.pep.*
11: /SIDS5/ptcodata/1/pubppaa/US11_NEW_PUB.pep1.*
12: /SIDS5/ptcodata/1/pubppaa/US60_NEW_PUB.pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	48	100.0	151	9	US-10-530-253-13
2	48	100.0	158	11	US-11-206-138-3
3	48	100.0	248	9	US-10-530-253-1
4	48	100.0	248	9	US-10-530-253-3
5	48	100.0	248	9	US-10-530-253-5
6	48	100.0	248	9	US-10-530-253-7
7	48	100.0	248	9	US-10-530-253-9
8	48	100.0	248	9	US-10-530-253-11
9	48	100.0	256	11	US-11-192-923A-2
10	43	89.6	158	9	US-10-530-253-15
11	43	89.6	158	9	US-10-530-253-20
12	37	77.1	158	9	US-10-530-253-19
13	35	72.9	52	11	US-11-096-568A-1159
14	35	72.9	615	11	US-11-188-298-4756
15	34	70.8	307	9	US-10-055-877-62
16	34	70.8	307	9	US-10-055-877-232
17	34	70.8	707	9	US-10-963-439-6
18	34	70.8	728	10	US-11-311-555-18
19	34	70.8	728	10	US-11-311-561-18
20	34	70.8	728	11	US-11-072-512-3399
21	34	70.8	739	9	US-10-963-439-5

22	70.8	812	11	US-11-079-463-6485	Sequence 6485, Ap
23	68.8	175	11	US-11-188-298-4065	Sequence 4065, Ap
24	68.8	243	11	US-11-079-463-9899	Sequence 9899, Ap
25	68.8	394	11	US-11-072-512-3188	Sequence 3188, Ap
26	66.7	215	11	US-11-045-004-2686	Sequence 2686, Ap
27	66.7	345	11	US-11-124-368A-285	Sequence 285, Ap
28	66.7	473	9	US-10-203-486-10	Sequence 10, Appl
29	66.7	501	9	US-10-467-657-7682	Sequence 7682, Ap
30	66.7	553	11	US-11-103-957-61	Sequence 61, Appl
31	66.7	577	9	US-10-203-486-8	Sequence 8, Appl1
32	66.7	626	11	US-11-098-686-10124	Sequence 10124, A
33	66.7	686	11	US-11-079-463-6039	Sequence 6039, Ap
34	64.6	158	9	US-10-530-253-26	Sequence 26, Appl
35	64.6	197	11	US-11-079-463-7727	Sequence 7727, Ap
36	64.6	206	11	US-11-087-099-2315	Sequence 26047, A
37	64.6	226	11	US-11-096-568A-26045	Sequence 26045, A
38	64.6	229	9	US-10-242-586-8	Sequence 8, Appl1
39	64.6	229	9	US-10-242-586-8	Sequence 8, Appl1
40	64.6	229	9	US-10-243-116-8	Sequence 8, Appl1
41	64.6	229	9	US-10-243-136-8	Sequence 8, Appl1
42	64.6	229	9	US-10-243-189-8	Sequence 8, Appl1
43	64.6	229	9	US-10-243-215-8	Sequence 8, Appl1
44	64.6	229	9	US-10-243-236-8	Sequence 8, Appl1
45	64.6	229	9	US-10-243-298-8	Sequence 8, Appl1
46	64.6	229	9	US-10-243-304-8	Sequence 8, Appl1
47	64.6	229	9	US-10-243-318-8	Sequence 8, Appl1
48	64.6	229	9	US-10-243-345-8	Sequence 8, Appl1
49	64.6	229	9	US-10-243-357-8	Sequence 8, Appl1
50	64.6	229	9	US-10-245-083-8	Sequence 8, Appl1
51	64.6	229	9	US-10-247-015-8	Sequence 8, Appl1
52	64.6	229	11	US-11-072-512-2074	Sequence 2074, Ap
53	64.6	231	11	US-11-072-512-2202	Sequence 2202, Ap
54	64.6	231	11	US-11-096-568A-24700	Sequence 24700, A
55	64.6	248	11	US-11-087-099-7533	Sequence 7533, Ap
56	64.6	368	11	US-11-188-298-12221	Sequence 12221, A
57	64.6	368	11	US-11-188-298-20952	Sequence 20952, A
58	64.6	457	11	US-11-087-099-10361	Sequence 10361, A
59	64.6	616	9	US-10-467-657-1220	Sequence 1220, Ap
60	64.6	664	9	US-10-793-626-1258	Sequence 1258, Ap
61	64.6	752	11	US-11-188-298-18269	Sequence 18269, A
62	64.6	974	9	US-10-995-561-895	Sequence 895, Ap
63	64.6	1098	11	US-11-072-512-2475	Sequence 2475, Ap
64	64.6	1223	9	US-10-506-444-1596	Sequence 1596, Ap
65	62.5	144	11	US-11-124-367A-276	Sequence 276, Ap
66	62.5	172	11	US-11-188-298-17891	Sequence 17891, A
67	62.5	186	11	US-11-188-298-4956	Sequence 4956, Ap
68	62.5	187	11	US-11-087-099-9124	Sequence 9124, Ap
69	62.5	201	11	US-11-096-568A-2939	Sequence 2939, Ap
70	62.5	231	11	US-11-096-568A-2938	Sequence 2938, Ap
71	62.5	258	7	US-09-995-493-216	Sequence 216, Ap
72	62.5	276	11	US-11-087-099-10298	Sequence 10298, A
73	62.5	287	11	US-11-096-568A-2937	Sequence 2937, Ap
74	62.5	308	11	US-11-156-084-304	Sequence 304, Ap
75	62.5	308	11	US-11-098-686-11286	Sequence 11286, A
76	62.5	308	11	US-11-131-479-2	Sequence 2, Appl1
77	62.5	498	11	US-11-131-479-76	Sequence 76, Appl
78	62.5	522	8	US-10-505-928-241	Sequence 241, Ap
79	62.5	522	11	US-11-131-479-7	Sequence 7, Appl1
80	62.5	522	11	US-11-131-479-9	Sequence 9, Appl1
81	62.5	567	9	US-10-420-192-8	Sequence 8, Appl1
82	62.5	567	11	US-11-072-175-197	Sequence 197, Ap
83	62.5	569	11	US-11-205-109-40	Sequence 40, Appl
84	62.5	877	9	US-10-242-586-90	Sequence 90, Appl
85	62.5	877	9	US-10-242-586-90	Sequence 90, Appl
86	62.5	877	9	US-10-243-116-90	Sequence 90, Appl
87	62.5	877	9	US-10-243-136-90	Sequence 90, Appl
88	62.5	877	9	US-10-243-189-90	Sequence 90, Appl
89	62.5	877	9	US-10-243-215-90	Sequence 90, Appl
90	62.5	877	9	US-10-243-236-90	Sequence 90, Appl
91	62.5	877	9	US-10-243-298-90	Sequence 90, Appl
92	62.5	877	9	US-10-243-304-90	Sequence 90, Appl
93	62.5	877	9	US-10-243-318-90	Sequence 90, Appl
94	62.5	877	9	US-10-243-345-90	Sequence 90, Appl

95	30	62.5	877	9	US-10-243-345-90	Sequence 90, Appl	168	29	60.4	520	9	US-10-137-873-144	Sequence 144, App
96	30	62.5	877	9	US-10-243-357-90	Sequence 90, Appl	169	29	60.4	520	9	US-10-152-370-144	Sequence 144, App
97	30	62.5	877	9	US-10-245-083-90	Sequence 90, Appl	170	29	60.4	520	11	US-11-290-153-14	Sequence 144, App
98	30	62.5	877	9	US-10-247-015-90	Sequence 90, Appl	171	29	60.4	524	11	US-11-118-809-4	Sequence 4, Appl1
99	30	62.5	1032	11	US-11-014-367-3	Sequence 3, Appl1	172	29	60.4	528	11	US-11-118-809-2	Sequence 2, Appl1
100	30	62.5	1051	9	US-10-204-639-15	Sequence 15, Appl	173	29	60.4	546	9	US-10-194-487-412	Sequence 412, App
101	30	62.5	1767	9	US-10-204-639-54	Sequence 54, Appl	174	29	60.4	546	9	US-10-195-883-412	Sequence 412, App
102	29.5	61.5	517	9	US-10-641-678-47	Sequence 47, Appl	175	29	60.4	546	9	US-10-195-888-412	Sequence 412, App
103	29	60.4	18	11	US-11-004-399-2919	Sequence 2919, Ap	176	29	60.4	546	9	US-10-195-889-412	Sequence 412, App
104	29	60.4	50	11	US-11-144-947-425	Sequence 425, App	177	29	60.4	546	9	US-10-218-784-124	Sequence 124, App
105	29	60.4	70	11	US-11-144-947-581	Sequence 581, App	178	29	60.4	546	9	US-10-219-061-124	Sequence 124, App
106	29	60.4	75	11	US-11-096-568-770	Sequence 770, App	179	29	60.4	546	9	US-10-219-062-124	Sequence 124, App
107	29	60.4	93	11	US-11-188-298-16085	Sequence 16085, A	180	29	60.4	546	9	US-10-219-064-124	Sequence 124, App
108	29	60.4	106	11	US-11-264-096-791	Sequence 791, App	181	29	60.4	546	9	US-10-233-134-124	Sequence 124, App
109	29	60.4	135	11	US-11-079-463-6954	Sequence 6954, Ap	182	29	60.4	546	11	US-11-124-367A-198	Sequence 498, App
110	29	60.4	156	11	US-11-072-512-2001	Sequence 2001, Ap	183	29	60.4	546	11	US-11-124-367A-199	Sequence 499, App
111	29	60.4	207	11	US-11-096-568-33943	Sequence 33943, A	184	29	60.4	590	9	US-10-330-773-124	Sequence 124, App
112	29	60.4	222	11	US-11-096-568-33942	Sequence 33942, A	185	29	60.4	594	9	US-10-330-773-121	Sequence 121, App
113	29	60.4	228	11	US-11-264-096-492	Sequence 492, App	186	29	60.4	602	11	US-11-188-298-20838	Sequence 20838 A
114	29	60.4	232	11	US-11-096-568-33941	Sequence 33941, A	187	29	60.4	605	9	US-10-821-234-1207	Sequence 1207, Ap
115	29	60.4	267	11	US-11-140-024-5	Sequence 5, Appl1	188	29	60.4	660	9	US-10-469-469-139	Sequence 139, App
116	29	60.4	274	11	US-11-096-568-22762	Sequence 22762, A	189	29	60.4	660	11	US-11-079-463-6259	Sequence 6259, Ap
117	29	60.4	281	11	US-11-172-740-1248	Sequence 1248, Ap	190	29	60.4	786	11	US-11-072-512-2944	Sequence 2944, Ap
118	29	60.4	293	11	US-11-079-463-6530	Sequence 6530, Ap	191	29	60.4	798	11	US-11-031-206-192	Sequence 192, App
119	29	60.4	324	11	US-11-188-298-16407	Sequence 16407, A	192	29	60.4	891	11	US-11-182-016-38	Sequence 38, Appl
120	29	60.4	330	11	US-11-072-512-2839	Sequence 2839, A	193	29	60.4	941	9	US-10-501-035-341	Sequence 343, App
121	29	60.4	340	11	US-11-096-568-19051	Sequence 19051, A	194	29	60.4	1022	11	US-11-087-039-7499	Sequence 7499, Ap
122	29	60.4	341	9	US-10-793-626-3248	Sequence 3248, A	195	29	60.4	1042	11	US-11-037-243-74	Sequence 74, Appl
123	29	60.4	342	9	US-10-491-468-8	Sequence 8, Appl1	196	29	60.4	1249	9	US-10-506-454-486	Sequence 486, App
124	29	60.4	367	11	US-11-188-298-15884	Sequence 15884, A	197	29	60.4	1822	8	US-10-505-928-700	Sequence 700, App
125	29	60.4	370	11	US-11-188-298-12247	Sequence 12247, A	198	29	60.4	1822	11	US-11-169-041-193	Sequence 193, App
126	29	60.4	380	11	US-11-087-009-6102	Sequence 6102, Ap	199	29	60.4	1968	9	US-10-915-002-314	Sequence 314, App
127	29	60.4	380	11	US-11-188-298-5571	Sequence 5571, Ap	200	29	60.4	3390	9	US-10-204-252-20	Sequence 20, Appl
128	29	60.4	383	11	US-11-144-947-460	Sequence 460, App	201	29	60.4	3390	9	US-10-204-252-20	Sequence 22, Appl
129	29	60.4	387	9	US-10-506-454-865	Sequence 865, App	202	29	58.3	15	11	US-11-144-947-510	Sequence 510, App
130	29	60.4	408	9	US-10-491-468-13	Sequence 13, Appl	203	28	58.3	111	11	US-11-072-512-3831	Sequence 3831, Ap
131	29	60.4	408	11	US-11-072-512-2081	Sequence 2081, Ap	204	28	58.3	115	9	US-10-519-390-16	Sequence 16, Appl
132	29	60.4	429	11	US-11-050-857-637	Sequence 637, App	205	28	58.3	115	11	US-11-176-830-208	Sequence 208, App
133	29	60.4	429	11	US-11-051-720-1317	Sequence 1317, Ap	206	28	58.3	115	11	US-11-176-830-558	Sequence 568, App
134	29	60.4	432	9	US-10-878-556A-72	Sequence 72, Appl	207	28	58.3	115	11	US-11-176-830-558	Sequence 569, App
135	29	60.4	434	11	US-11-050-857-635	Sequence 635, App	208	28	58.3	115	11	US-11-176-830-570	Sequence 570, App
136	29	60.4	444	11	US-11-051-720-1315	Sequence 1315, Ap	209	28	58.3	115	11	US-11-176-830-572	Sequence 572, App
137	29	60.4	446	11	US-11-050-857-633	Sequence 633, App	210	28	58.3	115	11	US-11-176-830-573	Sequence 573, App
138	29	60.4	446	11	US-11-051-720-1430	Sequence 1430, Ap	211	28	58.3	115	11	US-11-176-830-577	Sequence 577, App
139	29	60.4	448	11	US-11-079-463-5912	Sequence 5912, Ap	212	28	58.3	115	11	US-11-176-830-578	Sequence 578, App
140	29	60.4	454	11	US-11-050-857-636	Sequence 636, App	213	28	58.3	115	11	US-11-176-830-578	Sequence 579, App
141	29	60.4	454	11	US-11-051-720-1316	Sequence 1316, Ap	214	28	58.3	115	11	US-11-176-830-580	Sequence 580, App
142	29	60.4	459	11	US-11-096-568-13083	Sequence 13083, A	215	28	58.3	115	11	US-11-176-830-581	Sequence 581, App
143	29	60.4	462	11	US-11-124-367A-497	Sequence 497, App	216	28	58.3	115	11	US-11-176-830-582	Sequence 582, App
144	29	60.4	464	11	US-11-050-857-634	Sequence 634, App	217	28	58.3	115	11	US-11-176-830-582	Sequence 583, App
145	29	60.4	464	11	US-11-051-720-1314	Sequence 1314, Ap	218	28	58.3	115	11	US-11-176-830-583	Sequence 584, App
146	29	60.4	469	11	US-11-188-298-8536	Sequence 8536, Ap	219	28	58.3	115	11	US-11-176-830-585	Sequence 585, App
147	29	60.4	469	11	US-11-188-298-8688	Sequence 8688, Ap	220	28	58.3	115	11	US-11-176-830-585	Sequence 586, App
148	29	60.4	469	11	US-11-188-298-19251	Sequence 19251, A	221	28	58.3	115	11	US-11-176-830-586	Sequence 587, App
149	29	60.4	473	11	US-11-188-298-21659	Sequence 21659, A	222	28	58.3	115	11	US-11-176-830-588	Sequence 588, App
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151	29	60.4	485	11	US-11-096-568A-13081	Sequence 13081, A	224	28	58.3	115	11	US-11-176-830-589	Sequence 590, App
152	29	60.4	504	11	US-11-188-298-2121	Sequence 2121, Ap	225	28	58.3	115	11	US-11-176-830-591	Sequence 591, App
153	29	60.4	504	11	US-11-188-298-17474	Sequence 17474, A	226	28	58.3	115	11	US-11-176-830-592	Sequence 592, App
154	29	60.4	504	11	US-11-188-298-22138	Sequence 22138, A	227	28	58.3	115	11	US-11-176-830-593	Sequence 593, App
155	29	60.4	513	11	US-11-087-099-10936	Sequence 10936, Ap	228	28	58.3	115	11	US-11-176-830-594	Sequence 594, App
156	29	60.4	513	11	US-11-188-298-12045	Sequence 12045, A	229	28	58.3	115	11	US-11-176-830-595	Sequence 595, App
157	29	60.4	518	11	US-11-024-959-397	Sequence 397, App	230	28	58.3	115	11	US-11-176-830-595	Sequence 596, App
158	29	60.4	519	11	US-11-145-4058-26	Sequence 26, Appl	231	28	58.3	115	11	US-11-176-830-596	Sequence 597, App
159	29	60.4	519	11	US-11-145-4058-28	Sequence 28, Appl	232	28	58.3	115	11	US-11-176-830-597	Sequence 598, App
160	29	60.4	519	11	US-11-145-4058-30	Sequence 30, Appl	233	28	58.3	115	11	US-11-176-830-598	Sequence 599, App
161	29	60.4	519	11	US-11-145-4058-32	Sequence 32, Appl	234	28	58.3	115	11	US-11-176-830-600	Sequence 600, App
162	29	60.4	519	11	US-11-145-4058-34	Sequence 34, Appl	235	28	58.3	115	11	US-11-176-830-601	Sequence 601, App
163	29	60.4	519	11	US-11-145-4058-36	Sequence 36, Appl	236	28	58.3	115	11	US-11-176-830-601	Sequence 602, App
164	29	60.4	519	11	US-11-145-4058-38	Sequence 38, Appl	237	28	58.3	126	11	US-11-096-568A-21236	Sequence 21236, A
165	29	60.4	520	9	US-10-131-826A-144	Sequence 144, App	238	28	58.3	128	9	US-10-475-075-198	Sequence 198, App
166	29	60.4	520	9	US-10-973-115B-144	Sequence 144, App	239	28	58.3	128	9	US-10-475-075-198	Sequence 198, App
167	29	60.4	520	9	US-10-973-115B-144	Sequence 144, App	240	28	58.3	128	9	US-10-475-075-198	Sequence 144, App

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242	28	58.3	130	9	US-10-467-657-7656	Sequence 7656, Ap	315	28	58.3	437	11	US-11-188-298-5230	Sequence 5230, Ap
243	28	58.3	130	9	US-10-467-657-8416	Sequence 8416, Ap	316	28	58.3	445	11	US-11-096-568A-33243	Sequence 33243, Ap
244	28	58.3	166	11	US-11-094-519A-47	Sequence 47, Appl	317	28	58.3	450	11	US-11-024-959-396	Sequence 396, App
245	28	58.3	172	11	US-11-177-010-47	Sequence 2, Appl1	318	28	58.3	450	11	US-11-079-463-8683	Sequence 8683, Ap
246	28	58.3	172	11	US-11-177-010-4	Sequence 4, Appl1	319	28	58.3	456	11	US-11-188-298-2251	Sequence 2251, Ap
247	28	58.3	172	11	US-11-112-369-1	Sequence 1, Appl1	320	28	58.3	457	9	US-10-063-703-12	Sequence 12, Appl
248	28	58.3	172	11	US-11-112-369-2	Sequence 2, Appl1	321	28	58.3	457	9	US-10-194-487-48	Sequence 48, Appl
249	28	58.3	172	11	US-11-298-955-2	Sequence 2, Appl1	322	28	58.3	457	9	US-10-195-883-48	Sequence 48, Appl
250	28	58.3	172	11	US-11-298-955-3	Sequence 3, Appl1	323	28	58.3	457	9	US-10-195-889-48	Sequence 48, Appl
251	28	58.3	177	11	US-11-264-096-1721	Sequence 1719, Ap	324	28	58.3	457	9	US-10-216-161A-19	Sequence 19, Appl
252	28	58.3	177	11	US-11-264-096-1721	Sequence 1721, Ap	325	28	58.3	457	11	US-11-102-240-12	Sequence 12, Appl
253	28	58.3	178	7	US-09-995-493-82	Sequence 82, Appl	326	28	58.3	457	11	US-11-103-195-12	Sequence 12, Appl
254	28	58.3	203	11	US-11-079-463-9831	Sequence 9831, Ap	327	28	58.3	469	11	US-11-188-298-1906	Sequence 1906, Ap
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257	28	58.3	227	11	US-11-264-096-1722	Sequence 1722, Ap	330	28	58.3	471	11	US-11-188-298-5443	Sequence 5443, Ap
258	28	58.3	227	11	US-11-264-096-1722	Sequence 1723, Ap	331	28	58.3	482	11	US-11-188-298-6557	Sequence 6557, Ap
259	28	58.3	257	11	US-11-096-568A-9050	Sequence 9050, Ap	332	28	58.3	497	11	US-11-188-298-6557	Sequence 17543, A
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263	28	58.3	296	9	US-10-195-888-32	Sequence 32, Appl	336	28	58.3	502	11	US-11-188-298-2892	Sequence 7155, Ap
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266	28	58.3	308	11	US-11-188-298-9050	Sequence 9050, Ap	339	28	58.3	508	11	US-11-188-298-17885	Sequence 17685, A
267	28	58.3	313	11	US-11-087-099-9808	Sequence 9808, Ap	340	28	58.3	509	11	US-11-188-298-17118	Sequence 17118, A
268	28	58.3	314	11	US-11-098-686-10957	Sequence 10957, A	341	28	58.3	512	11	US-11-188-298-17118	Sequence 500, App
269	28	58.3	317	11	US-11-094-519A-38	Sequence 38, Appl	342	28	58.3	514	11	US-11-024-959-500	Sequence 1066, Ap
270	28	58.3	319	9	US-10-511-538-21	Sequence 21, Appl	343	28	58.3	522	11	US-11-045-004-1066	Sequence 38, Appl
271	28	58.3	322	11	US-11-067-121-15	Sequence 15, Appl	344	28	58.3	551	9	US-10-491-468-18	Sequence 33241, A
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273	28	58.3	323	9	US-10-931-198-58	Sequence 58, Appl	346	28	58.3	602	11	US-11-131-263-19	Sequence 19, Appl
274	28	58.3	323	9	US-10-942-042-58	Sequence 58, Appl	347	28	58.3	602	11	US-11-131-263-29	Sequence 29, Appl
275	28	58.3	326	11	US-11-072-512-3428	Sequence 3428, Ap	348	28	58.3	610	11	US-11-079-463-9122	Sequence 9122, Ap
276	28	58.3	343	9	US-11-096-568A-6730	Sequence 6730, Ap	349	28	58.3	643	9	US-10-504-544-2	Sequence 2, Appl1
277	28	58.3	347	9	US-10-516-635-6	Sequence 6, Appl1	350	28	58.3	660	11	US-11-188-298-17672	Sequence 17672, A
278	28	58.3	348	9	US-11-067-121-5	Sequence 4, Appl1	351	28	58.3	660	11	US-10-055-877-921	Sequence 321, App
279	28	58.3	349	11	US-11-067-121-5	Sequence 56, Appl1	352	28	58.3	693	9	US-11-189-301-20	Sequence 20, Appl
280	28	58.3	349	9	US-10-921-793-56	Sequence 56, Appl	353	28	58.3	701	11	US-11-189-301-19	Sequence 19, Appl
281	28	58.3	349	9	US-10-931-198-56	Sequence 56, Appl	354	28	58.3	743	11	US-11-072-512-2340	Sequence 2340, Ap
282	28	58.3	349	9	US-10-942-042-56	Sequence 56, Appl	355	28	58.3	743	11	US-11-072-512-2340	Sequence 30499, A
283	28	58.3	349	9	US-10-516-635-2	Sequence 2, Appl1	356	28	58.3	750	11	US-11-189-301-21	Sequence 21, Appl
284	28	58.3	352	11	US-11-067-121-14	Sequence 14, Appl	357	28	58.3	766	11	US-11-096-568A-30499	Sequence 10359, A
285	28	58.3	352	11	US-11-096-568A-6729	Sequence 6729, Ap	358	28	58.3	787	11	US-11-087-099-10359	Sequence 10681, A
286	28	58.3	358	9	US-10-055-877-176	Sequence 1766, App	359	28	58.3	787	11	US-11-087-099-110815	Sequence 11835, A
287	28	58.3	358	11	US-11-174-816-64	Sequence 69, Appl	360	28	58.3	787	11	US-11-188-298-20849	Sequence 20849, A
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290	28	58.3	358	11	US-11-174-819-83	Sequence 88, Appl	363	28	58.3	788	11	US-11-087-099-7244	Sequence 7520, Ap
291	28	58.3	358	11	US-11-174-819-89	Sequence 278, App	364	28	58.3	788	11	US-11-087-099-7520	Sequence 9550, Ap
292	28	58.3	359	11	US-11-188-298-778	Sequence 70, Appl	365	28	58.3	788	11	US-11-188-298-6897	Sequence 6897, Ap
293	28	58.3	362	11	US-11-174-816-70	Sequence 9061, Ap	366	28	58.3	788	11	US-11-188-298-17710	Sequence 17710, A
294	28	58.3	365	11	US-11-188-298-9081	Sequence 39, Appl	367	28	58.3	792	11	US-11-088-686-29	Sequence 29, Appl
295	28	58.3	366	8	US-10-505-928-39	Sequence 39, Appl	368	28	58.3	870	11	US-11-087-099-1636	Sequence 1636, Ap
296	28	58.3	367	11	US-11-096-568A-21787	Sequence 21787, Ap	369	28	58.3	870	11	US-11-087-099-9223	Sequence 9223, Ap
297	28	58.3	374	9	US-10-793-625-3096	Sequence 3096, Ap	370	28	58.3	870	11	US-11-188-298-8577	Sequence 8577, Ap
298	28	58.3	378	11	US-11-098-686-10740	Sequence 10740, A	371	28	58.3	870	11	US-11-188-298-12564	Sequence 12564, A
299	28	58.3	380	11	US-11-188-298-20598	Sequence 20598, A	372	28	58.3	885	11	US-11-096-568A-30498	Sequence 30498, A
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309	28	58.3	414	11	US-11-188-298-2117	Sequence 2117, Ap	375	28	58.3	934	11	US-11-024-96-553-4	Sequence 4, Appl1
310	28	58.3	415	11	US-11-096-568A-21786	Sequence 21786, A	376	28	58.3	943	11	US-11-124-168A-293	Sequence 293, App
311	28	58.3	422	9	US-10-506-454-1401	Sequence 1401, Ap	377	28	58.3	943	11	US-11-124-168A-295	Sequence 146, App
312	28	58.3	422	9	US-11-096-568A-24469	Sequence 24469, A	378	28	58.3	945	9	US-10-131-826A-146	Sequence 146, App
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							380	28	58.3	945	9	US-10-137-873A-146	Sequence 146, App
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							382	28	58.3	945	9	US-10-137-873A-146	Sequence 146, App
							383	28	58.3	945	11	US-11-019-711-38	Sequence 38, Appl
							384	28	58.3	945	11	US-11-019-711-38	Sequence 38, Appl
							385	28	58.3	945	11	US-11-183-136-18	Sequence 18, Appl
							386	28	58.3	945	11	US-11-183-136-20	Sequence 20, Appl

387	28	58.3	945	11	US-11-290-153-146	Sequence 146, App	460	27	56.2	148	11	US-11-096-568A-29967	Sequence 29967, A
388	28	58.3	1047	11	US-11-072-512-2408	Sequence 2408, Ap	461	27	56.2	150	9	US-10-469-561-25	Sequence 25, Appl
389	28	58.3	1342	9	US-10-770-726-63	Sequence 63, Appl	462	27	56.2	152	11	US-11-079-463-6202	Sequence 4620, Ap
390	28	58.3	1342	11	US-11-113-202-12	Sequence 12, Appl	463	27	56.2	153	8	US-10-511-937-2476	Sequence 2476, Ap
391	28	58.3	1342	11	US-11-113-202-14	Sequence 14, Appl	464	27	56.2	153	11	US-11-174-398-8	Sequence 8, Appl1
392	28	58.3	1450	11	US-11-019-711-113	Sequence 113, Appl	465	27	56.2	153	11	US-11-289-226-13	Sequence 13, Appl1
393	28	58.3	1474	9	US-10-995-561-873	Sequence 873, App	466	27	56.2	158	8	US-10-505-928-441	Sequence 2983, Ap
394	28	58.3	1474	9	US-10-877-346-13	Sequence 13, Appl	467	27	56.2	158	8	US-10-511-937-2993	Sequence 2993, Ap
395	28	58.3	1907	11	US-11-039-398-25	Sequence 25, Appl	468	27	56.2	158	9	US-10-995-561-1031	Sequence 1031, Ap
396	28	58.3	3387	9	US-10-204-252-24	Sequence 24, Appl	469	27	56.2	158	9	US-10-995-561-1032	Sequence 1032, Ap
397	28	58.3	3387	9	US-10-204-252-26	Sequence 26, Appl	470	27	56.2	158	9	US-10-784-004-664	Sequence 664, App
398	28	58.3	3389	9	US-10-204-252-10	Sequence 10, Appl	471	27	56.2	160	9	US-10-530-253-25	Sequence 25, Appl
399	28	58.3	3391	9	US-10-204-252-6	Sequence 6, Appl1	472	27	56.2	160	11	US-11-210-756-15	Sequence 15, Appl
400	28	58.3	3391	9	US-10-204-252-8	Sequence 8, Appl1	473	27	56.2	165	11	US-11-175-690-99	Sequence 99, Appl
401	28	58.3	3391	9	US-10-204-252-12	Sequence 12, Appl	474	27	56.2	165	11	US-11-175-690-517	Sequence 517, App
402	28	58.3	3391	9	US-10-204-252-14	Sequence 14, Appl	475	27	56.2	166	11	US-11-133-722-45	Sequence 45, Appl
403	28	58.3	3391	9	US-10-204-252-16	Sequence 16, Appl	476	27	56.2	166	11	US-11-176-830-186	Sequence 186, App
404	28	58.3	3391	9	US-10-204-252-28	Sequence 28, Appl	477	27	56.2	166	11	US-11-177-409-2	Sequence 2, Appl1
405	28	58.3	3402	9	US-10-204-252-18	Sequence 18, Appl	478	27	56.2	166	11	US-11-198-765-83	Sequence 83, Appl
406	28	58.3	3623	9	US-10-995-561-593	Sequence 593, App	479	27	56.2	167	11	US-11-175-690-103	Sequence 103, App
407	27	56.2	8	11	US-11-045-024-1620	Sequence 1620, App	480	27	56.2	168	9	US-10-454-437-196	Sequence 196, App
408	27	56.2	8	11	US-11-045-024-4333	Sequence 4333, Ap	481	27	56.2	168	11	US-11-055-823-360	Sequence 360, App
409	27	56.2	8	11	US-11-045-024-7592	Sequence 7592, Ap	482	27	56.2	170	11	US-11-096-568A-29966	Sequence 29966, A
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411	27	56.2	9	11	US-11-045-024-4445	Sequence 4445, Ap	484	27	56.2	175	11	US-11-087-099-9020	Sequence 9020, Ap
412	27	56.2	9	11	US-11-045-024-7595	Sequence 7595, Ap	485	27	56.2	185	9	US-10-467-657-1662	Sequence 1662, Ap
413	27	56.2	9	11	US-11-045-024-13908	Sequence 13908, A	486	27	56.2	189	11	US-11-147-492-6	Sequence 6, Appl
414	27	56.2	9	11	US-11-045-024-14183	Sequence 14183, A	487	27	56.2	189	11	US-11-147-492-22	Sequence 22, Appl1
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417	27	56.2	10	11	US-11-045-024-2067	Sequence 2067, Ap	490	27	56.2	207	11	US-11-126-427-6	Sequence 6, Appl1
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419	27	56.2	10	11	US-11-045-024-6582	Sequence 6582, Ap	492	27	56.2	210	7	US-09-978-3604-432	Sequence 432, App
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430	27	56.2	80	11	US-11-096-568A-2785	Sequence 2785, Ap	503	27	56.2	243	9	US-10-973-1189-362	Sequence 362, App
431	27	56.2	83	9	US-10-467-657-7970	Sequence 7970, Ap	504	27	56.2	243	9	US-10-137-8734-362	Sequence 362, App
432	27	56.2	99	9	US-10-821-234-1702	Sequence 1702, Ap	505	27	56.2	243	9	US-10-152-370-362	Sequence 362, App
433	27	56.2	102	11	US-11-096-568A-8713	Sequence 8713, Ap	506	27	56.2	244	11	US-11-290-153-362	Sequence 362, App
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446	27	56.2	129	11	US-11-176-830-555	Sequence 555, App	519	27	56.2	287	11	US-11-188-299-3945	Sequence 3945, Ap
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455	27	56.2	129	11	US-11-176-830-564	Sequence 564, App	528	27	56.2	325	9	US-10-973-1185-516	Sequence 516, App
456	27	56.2	129	11	US-11-176-830-565	Sequence 565, App	529	27	56.2	325	9	US-10-213-535-11	Sequence 11, Appl
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458	27	56.2	129	11	US-11-176-830-567	Sequence 567, App	531	27	56.2	325	9	US-10-152-370-516	Sequence 516, App
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550	27	56.2	426	9	US-10-455-772-158	Sequence 158, App	623	27	56.2	735	9	US-10-194-487-88	Sequence 88, Appl
551	27	56.2	444	11	US-11-096-568A-11404	Sequence 11404, A	624	27	56.2	735	9	US-10-195-883-88	Sequence 88, Appl
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556	27	56.2	453	9	US-10-131-826A-270	Sequence 270, App	629	27	56.2	739	11	US-11-009-063-31	Sequence 31, Appl
557	27	56.2	453	9	US-10-973-115B-270	Sequence 270, App	630	27	56.2	741	11	US-11-188-298-80A0	Sequence 80A0, Ap
558	27	56.2	453	9	US-10-137-873A-270	Sequence 270, App	631	27	56.2	747	9	US-10-131-826A-426	Sequence 426, App
559	27	56.2	453	9	US-10-153-370-270	Sequence 270, App	632	27	56.2	747	9	US-10-973-115B-426	Sequence 426, App
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562	27	56.2	456	11	US-10-821-234-1185	Sequence 1185, Ap	635	27	56.2	747	9	US-10-152-370-426	Sequence 426, App
563	27	56.2	457	11	US-11-079-463-6587	Sequence 6587, Ap	636	27	56.2	747	11	US-11-390-153-426	Sequence 426, App
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568	27	56.2	467	11	US-11-188-298-16761	Sequence 16761, A	641	27	56.2	787	9	US-10-455-772-152	Sequence 152, App
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584	27	56.2	503	11	US-11-188-298-14816	Sequence 14816, A	657	27	56.2	802	9	US-10-245-083-78	Sequence 78, Appl
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589	27	56.2	523	11	US-11-045-004-505	Sequence 505, App	662	27	56.2	813	9	US-10-194-487-466	Sequence 466, App
590	27	56.2	525	9	US-10-641-678-52	Sequence 52, App	663	27	56.2	813	9	US-10-195-883-466	Sequence 466, App
591	27	56.2	527	11	US-11-096-568A-28901	Sequence 28901, A	664	27	56.2	813	9	US-10-195-888-466	Sequence 466, App
592	27	56.2	530	11	US-11-096-568A-15782	Sequence 15782, A	665	27	56.2	813	9	US-10-195-889-466	Sequence 466, App
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596	27	56.2	541	11	US-11-010-239-95	Sequence 95, Appl	669	27	56.2	830	11	US-11-096-568A-27818	Sequence 27818, A
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598	27	56.2	545	11	US-10-853-807A-9	Sequence 9, Appl1	671	27	56.2	832	11	US-11-031-006-188	Sequence 188, App
599	27	56.2	561	11	US-11-029-465-12	Sequence 12, Appl	672	27	56.2	870	9	US-10-455-772-162	Sequence 162, App
600	27	56.2	562	9	US-10-507-928-10	Sequence 10, Appl	673	27	56.2	874	9	US-10-455-772-160	Sequence 160, App
601	27	56.2	562	9	US-10-507-928-12	Sequence 12, Appl	674	27	56.2	895	9	US-11-079-463-9074	Sequence 9074, Ap
602	27	56.2	562	11	US-11-029-465-10	Sequence 10, Appl	675	27	56.2	905	11	US-11-186-284-2	Sequence 2, Appl1
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681	27	56.2	960	11	US-11-169-041-177	Sequence 177, App	754	26	54.2	209	9	US-10-195-889-306	Sequence 306, App
682	27	56.2	1001	11	US-11-132-285-40	Sequence 40, App	755	26	54.2	209	9	US-10-218-784-130	Sequence 100, App
683	27	56.2	1003	11	US-11-204-755-7	Sequence 7, App	756	26	54.2	209	9	US-10-219-061-100	Sequence 100, App
684	27	56.2	1003	11	US-11-204-755-9	Sequence 9, App	757	26	54.2	209	9	US-10-219-062-100	Sequence 100, App
685	27	56.2	1003	11	US-11-204-755-11	Sequence 11, App	758	26	54.2	209	9	US-10-219-064-100	Sequence 100, App
686	27	56.2	1013	9	US-10-131-826A-38	Sequence 38, App	759	26	54.2	209	9	US-10-233-134-100	Sequence 100, App
687	27	56.2	1013	9	US-10-973-115B-38	Sequence 38, App	760	26	54.2	209	11	US-11-102-240-96	Sequence 96, App
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689	27	56.2	1013	9	US-10-152-370-38	Sequence 38, App	762	26	54.2	209	11	US-11-103-195-96	Sequence 96, App
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973 26 54.2 708 9 US-10-973-115B-298 Sequence 298, App
974 26 54.2 708 9 US-10-137-873A-298 Sequence 298, App
975 26 54.2 708 9 US-10-152-370-298 Sequence 298, App
976 26 54.2 708 11 US-11-290-153-298 Sequence 298, App
977 26 54.2 709 11 US-11-188-298-17090 Sequence 17090, A
978 26 54.2 719 9 US-10-467-657-762 Sequence 762, App
979 26 54.2 720 11 US-11-113-424-28 Sequence 28, App
980 26 54.2 721 11 US-11-087-099-8786 Sequence 8786, App
981 26 54.2 724 11 US-11-096-568A-11882 Sequence 11882, A
982 26 54.2 725 11 US-11-087-099-4908 Sequence 4908, Ap
983 26 735 11 US-11-188-298-15331 Sequence 15331, A
984 26 54.2 747 11 US-11-113-424-26 Sequence 26, App
985 26 54.2 747 11 US-11-264-096-774 Sequence 774, App
986 26 54.2 749 11 US-11-098-686-10505 Sequence 10505, A
987 26 54.2 754 9 US-10-467-962B-63 Sequence 63, Appl
988 26 54.2 757 11 US-11-031-206-184 Sequence 184, App
989 26 54.2 758 11 US-11-188-298-21656 Sequence 21656, A
990 26 54.2 761 11 US-11-057-047-6 Sequence 6, Appl
991 26 54.2 762 9 US-10-506-454-648 Sequence 648, App
992 26 54.2 776 9 US-10-821-234-1171 Sequence 1171, Ap
993 26 54.2 791 11 US-11-087-099-7833 Sequence 7833, Ap
994 26 54.2 811 8 US-10-511-937-2584 Sequence 2584, Ap
995 26 54.2 816 11 US-11-072-512-2341 Sequence 2341, Ap
996 26 54.2 830 9 US-10-921-793-38 Sequence 38, Appl
997 26 54.2 830 9 US-10-931-198-38 Sequence 38, Appl
998 26 54.2 830 9 US-10-942-042-38 Sequence 38, Appl
999 26 54.2 835 11 US-11-098-686-10397 Sequence 10397, A
1000 26 54.2 836 8 US-10-511-937-2988 Sequence 2988, Ap
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ALIGNMENTS

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RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13
Query Match 100.0%; Score 48; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1
```

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; GENERAL INFORMATION:
; APPLICANT: Healthbanc Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; CURRENT FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3
```

```
Query Match 100.0%; Score 48; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.041;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-1
Query Match 100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
```

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; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
; US-10-530-253-7
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 3
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-3
```

```
Query Match          100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPPQCTEL 9
    |||||
Db 11 KLPPQCTEL 19
```

```
RESULT 5
US-10-530-253-5
; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5
```

```
Query Match          100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 KLPPQCTEL 9
    |||||
Db 11 KLPPQCTEL 19
```

```
RESULT 6
US-10-530-253-7
; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 7
; LENGTH: 248
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; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
; US-10-530-253-7
```

```
Query Match          100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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QY 1 KLPPQCTEL 9
    |||||
Db 108 KLPPQCTEL 116
```

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RESULT 7
US-10-530-253-9
; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9
```

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Query Match          100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
QY 1 KLPPQCTEL 9
    |||||
Db 108 KLPPQCTEL 116
```

```
RESULT 8
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11
```

```
Query Match          100.0%; Score 48; DB 9; Length 248;
```

Best Local Similarity 100.0%; Pred. No. 0.063;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 108 KLPOLCTEL 116

RESULT 9
US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018926A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; TITLE OF INVENTION: RECOMBINANT REPLICON
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; PRIOR FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 48; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.065;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 116 KLPOLCTEL 124

RESULT 10
US-10-530-253-15
; Sequence 15, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 15
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 18
US-10-530-253-15

Query Match 89.6%; Score 43; DB 9; Length 158;
Best Local Similarity 88.9%; Pred. No. 0.38;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 13 KLPOLCTEL 21

RESULT 11
US-10-530-253-20

; Sequence 20, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 45
US-10-530-253-20

Query Match 89.6%; Score 43; DB 9; Length 158;
Best Local Similarity 88.9%; Pred. No. 0.38;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 13 KLPOLCTEL 21

RESULT 12
US-10-530-253-19
; Sequence 19, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 19
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 39
US-10-530-253-19

Query Match 77.1%; Score 37; DB 9; Length 158;
Best Local Similarity 77.8%; Pred. No. 5.4;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 13 KLPOLCTEL 21

RESULT 13
US-11-096-568A-1159
; Sequence 1159, Application US/11096568A
; Publication No. US20060048240A1

```

; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Thereby
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1159
; LENGTH: 52
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(52)
; OTHER INFORMATION: Ceres Seq. ID no. 13600515
US-11-096-568A-1159

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Query Match 72.9%; Score 35; DB 11; Length 52;
Best Local Similarity 75.0%; Pred. No. 4.6;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

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QY 1 KLPOCTEL 8
DB 12 QLPQCTE 19

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RESULT 14
US-11-188-298-4756
; Sequence 4756, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 4756
; LENGTH: 615
; TYPE: PRT
; ORGANISM: Oryza sativa
US-11-188-298-4756

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```

Query Match 72.9%; Score 35; DB 11; Length 615;
Best Local Similarity 77.8%; Pred. No. 46;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

```

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QY 1 KLPOCTEL 9
DB 476 KLPELTEL 484

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RESULT 15
US-10-055-877-62
; Sequence 62, Application US/10055877
; Publication No. US20050288241A1
; GENERAL INFORMATION:
; APPLICANT: Decristofaro, Marc
; APPLICANT: Padigar, Muralidhara
; APPLICANT: Miller, Charles
; APPLICANT: Tchernev, Velizar
; APPLICANT: Zhong, Mei
; APPLICANT: Anderson, David
; APPLICANT: Ballinger, Robert
; APPLICANT: Gerlach, Valerie
; APPLICANT: Spytek, Kimberly
; APPLICANT: Ratelli, Luca
; APPLICANT: Kekuda, Ramesh
; APPLICANT: Guo, Xiaojia
; APPLICANT: Zehnusen, Bryan

```

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; APPLICANT: Andrew, David
; APPLICANT: Mezes, Peter
; APPLICANT: Patturajan, Meera
; APPLICANT: Burgess, Catherine
; APPLICANT: Eissen, Andrew
; APPLICANT: Wolenc, Adam
; APPLICANT: Baumgartner, Jason
; APPLICANT: Shinkets, Richard
; APPLICANT: Gusev, Vladimir
; APPLICANT: Verne, Corine
; APPLICANT: Taupier Jr., Raymond
; APPLICANT: Pena, Carol
; APPLICANT: Shenoy, Suresh
; APPLICANT: Li, Li
; APPLICANT: Casman, Stacie
; APPLICANT: Boldog, Ference
; TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
; FILE REFERENCE: 21402-251
; CURRENT APPLICATION NUMBER: US/10/055,877
; CURRENT FILING DATE: 2002-01-22
; PRIOR APPLICATION NUMBER: 60/262,892
; PRIOR FILING DATE: 2001-01-19
; PRIOR APPLICATION NUMBER: 60/263,598
; PRIOR FILING DATE: 2001-01-23
; PRIOR APPLICATION NUMBER: 60/263,799
; PRIOR FILING DATE: 2001-01-24
; PRIOR APPLICATION NUMBER: 60/264,117
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/264,139
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/264,478
; PRIOR FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: 60/263,351
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: 60/272,870
; PRIOR FILING DATE: 2001-03-02
; PRIOR APPLICATION NUMBER: 60/275,990
; PRIOR FILING DATE: 2001-03-14
; PRIOR APPLICATION NUMBER: 60/275,927
; PRIOR FILING DATE: 2001-03-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 512
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 62
; LENGTH: 307
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-62

```

```

Query Match 70.8%; Score 34; DB 9; Length 307;
Best Local Similarity 71.4%; Pred. No. 37;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 3 POLCTQM 142
DB 136 POLCTQM 142

```

```

RESULT 16
US-10-055-877-232
; Sequence 232, Application US/10055877
; Publication No. US20050288241A1
; GENERAL INFORMATION:
; APPLICANT: Decristofaro, Marc
; APPLICANT: Padigar, Muralidhara
; APPLICANT: Miller, Charles
; APPLICANT: Tchernev, Velizar
; APPLICANT: Zhong, Mei
; APPLICANT: Anderson, David
; APPLICANT: Ballinger, Robert
; APPLICANT: Gerlach, Valerie
; APPLICANT: Spytek, Kimberly
; APPLICANT: Ratelli, Luca

```

```

; APPLICANT: Kekuda, Ramesh
; APPLICANT: Guo, Xiaojia
; APPLICANT: Zerhusen, Bryan
; APPLICANT: Andrew, David
; APPLICANT: Mezes, Peter
; APPLICANT: Patturajan, Meera
; APPLICANT: Burgess, Catherine
; APPLICANT: Eileen, Andrew
; APPLICANT: Wolenc, Adam
; APPLICANT: Baumgartner, Jason
; APPLICANT: Shinkets, Richard
; APPLICANT: Gusev, Vladimir
; APPLICANT: Verter, Corine
; APPLICANT: Taupier Jr., Raymond
; APPLICANT: Pena, Carol
; APPLICANT: Shenoy, Suresh
; APPLICANT: Li, Li
; APPLICANT: Casman, Stacie
; APPLICANT: Boldog, Ference
; TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
; FILE REFERENCE: 21402-251
; CURRENT APPLICATION NUMBER: US/10/055,877
; CURRENT FILING DATE: 2002-01-22
; PRIOR APPLICATION NUMBER: 60/262,892
; PRIOR FILING DATE: 2001-01-19
; PRIOR APPLICATION NUMBER: 60/263,598
; PRIOR FILING DATE: 2001-01-23
; PRIOR APPLICATION NUMBER: 60/263,799
; PRIOR FILING DATE: 2001-01-24
; PRIOR APPLICATION NUMBER: 60/264,117
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/264,139
; PRIOR FILING DATE: 2001-01-25
; PRIOR APPLICATION NUMBER: 60/264,478
; PRIOR FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: 60/263,351
; PRIOR FILING DATE: 2001-01-30
; PRIOR APPLICATION NUMBER: 60/272,870
; PRIOR FILING DATE: 2001-03-02
; PRIOR APPLICATION NUMBER: 60/275,990
; PRIOR FILING DATE: 2001-03-14
; PRIOR APPLICATION NUMBER: 60/275,927
; PRIOR FILING DATE: 2001-03-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 512
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 232
; LENGTH: 307
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-055-877-232

Query Match          70.8%; Score 34; DB 9; Length 307;
Best Local Similarity 71.4%; Pred. No. 37;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy      3 POLCTEL 9
Db      136 POLCTQM 142

RESULT 17
US-10-963-439-6
; Sequence 6, Application US/10963439
; Publication No. US20060079444A1
; GENERAL INFORMATION:
; APPLICANT: Ron, Dina
; TITLE OF INVENTION: HUMAN SEF ISOFORMS AND METHODS OF USING SAME FOR CANCER GENE
; FILE REFERENCE: 28385
; CURRENT APPLICATION NUMBER: US/10/963,439
; CURRENT FILING DATE: 2004-10-11
; NUMBER OF SEQ ID NOS: 22
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; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 6
; LENGTH: 707
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-963-439-6

Query Match          70.8%; Score 34; DB 9; Length 707;
Best Local Similarity 75.0%; Pred. No. 81;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      2 LPOLCTEL 9
Db      472 LPOLCSHL 479

RESULT 18
US-11-311-555-18
; Sequence 18, Application US/11311555
; Publication No. US20060088916A1
; GENERAL INFORMATION:
; APPLICANT: Genentech, Inc.
; APPLICANT: Chen, Jian
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Rong, Sherman
; APPLICANT: Goddard, Audrey
; APPLICANT: Godowski, Paul
; APPLICANT: Grimaldi, Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Li, Hanzhong
; APPLICANT: Hillan, Kenneth
; APPLICANT: Tuma, Daniel
; APPLICANT: Vanlookeren, Menno
; APPLICANT: Vandlen, Richard
; APPLICANT: Watanabe, Colin
; APPLICANT: Williams, P. Mickey
; APPLICANT: Wood, William
; APPLICANT: Yansura, Daniel
; TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
; FILE REFERENCE: P1381R1C1P1(US)
; CURRENT APPLICATION NUMBER: US/11/311,555
; CURRENT FILING DATE: 2005-12-20
; PRIOR APPLICATION NUMBER: US/09/747,259
; PRIOR FILING DATE: 2000-12-20
; PRIOR APPLICATION NUMBER: US 09/311,832
; PRIOR FILING DATE: 1999-05-14
; PRIOR APPLICATION NUMBER: US 60/172,096
; PRIOR FILING DATE: 1999-12-23
; PRIOR APPLICATION NUMBER: PCT/US99/31274
; PRIOR FILING DATE: 1999-12-30
; PRIOR APPLICATION NUMBER: US 60/175,481
; PRIOR FILING DATE: 2000-01-11
; PRIOR APPLICATION NUMBER: PCT/US00/04341
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/US00/05841
; PRIOR FILING DATE: 2000-03-02
; PRIOR APPLICATION NUMBER: US 60/191,007
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: PCT/US00/07532
; PRIOR FILING DATE: 2000-03-21
; PRIOR APPLICATION NUMBER: PCT/US00/15264
; PRIOR FILING DATE: 2000-06-02
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 39
; SEQ ID NO 18
; LENGTH: 728
; TYPE: PRT
; ORGANISM: Homo Sapien
US-11-311-555-18

Query Match          70.8%; Score 34; DB 10; Length 728;
Best Local Similarity 75.0%; Pred. No. 83;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

Qy 2 LPOLCTEL 9
Db 493 LPOLCSHL 500

RESULT 19

US-11-311-561-18
; Sequence 18, Application US/11311561
; Publication No. US20060088917A1

GENERAL INFORMATION:

APPLICANT: Genentech, Inc.
APPLICANT: Chen, Jian
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Goddard, Audrey
APPLICANT: Godowski, Paul
APPLICANT: Grimaldi, Christopher
APPLICANT: Gurney, Auelin
APPLICANT: Li, Hanzhong
APPLICANT: Hillan, Kenneth
APPLICANT: Tumas, Daniel
APPLICANT: Vanlookeren, Menno
APPLICANT: Vandlen, Richard
APPLICANT: Watanabe, Colin
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William
APPLICANT: Yansura, Daniel
TITLE OF INVENTION: IL-17 HOMOLOGOUS POLYPEPTIDES AND THERAPEUTIC USES THEREOF
FILE REFERENCE: P1381R1C1P1 (US)
CURRENT APPLICATION NUMBER: US/11/311,561
CURRENT FILING DATE: 2005-12-20
PRIOR APPLICATION NUMBER: US/09/747,259
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: US 09/311,832
PRIOR FILING DATE: 1999-05-14
PRIOR APPLICATION NUMBER: US 60/172,096
PRIOR FILING DATE: 1999-12-23
PRIOR APPLICATION NUMBER: PCT/US99/31274
PRIOR FILING DATE: 1999-12-30
PRIOR APPLICATION NUMBER: US 60/175,481
PRIOR FILING DATE: 2000-01-11
PRIOR APPLICATION NUMBER: PCT/US00/04341
PRIOR FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/US00/05841
PRIOR FILING DATE: 2000-03-02
PRIOR APPLICATION NUMBER: US 60/191,007
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: PCT/US00/07532
PRIOR FILING DATE: 2000-03-21
PRIOR APPLICATION NUMBER: PCT/US00/15264
PRIOR FILING DATE: 2000-06-02
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 39
SEQ ID NO 18
LENGTH: 728
TYPE: PRT
ORGANISM: Homo Sapien
US-11-311-561-18

Query Match 70.8%; Score 34; DB 10; Length 728;
Best Local Similarity 75.0%; Pred. No. 83;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 LPOLCTEL 9
Db 493 LPOLCSHL 500

RESULT 20
US-11-072-512-3399
; Sequence 3399, Application US/11072512
; Publication No. US20060029945A1

GENERAL INFORMATION:

APPLICANT: ISOGAI, TAKAO
APPLICANT: SUGIYAMA, TOMOYASU
APPLICANT: OTSUKI, TETSUJI
APPLICANT: WAKAMATSU, AI
APPLICANT: SATO, HIROYUKI
APPLICANT: ISHII, SHIZUKO
APPLICANT: YAMAMOTO, JUN-ICHI
APPLICANT: ISONO, YUDKO
APPLICANT: HIO, YURI
APPLICANT: OTSUKA, KAORU
APPLICANT: NAGAI, KEIICHI
APPLICANT: IRIE, RYOTARO
APPLICANT: TAMECHIKA, ICHIRO
APPLICANT: SEKI, NAOHICO
APPLICANT: YOSHIKAWA, TSUTOMU
APPLICANT: OTSUKA, MOTYUKI
APPLICANT: NAGAHARI, KENJI
APPLICANT: MASUHO, YASUHIKO
TITLE OF INVENTION: Novel full length cDNA
FILE REFERENCE: 084335-0191
CURRENT APPLICATION NUMBER: US/11/072,512
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US 60/350,978
PRIOR FILING DATE: 2002-01-25
PRIOR APPLICATION NUMBER: JP 2001-379298
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 4096
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3399
LENGTH: 728
TYPE: PRT
ORGANISM: Homo sapiens
US-11-072-512-3399

Query Match 70.8%; Score 34; DB 11; Length 728;
Best Local Similarity 75.0%; Pred. No. 83;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 LPOLCTEL 9
Db 493 LPOLCSHL 500

RESULT 21

US-10-963-439-5
; Sequence 5, Application US/10963439
; Publication No. US20060079444A1

GENERAL INFORMATION:

APPLICANT: Kon, Dina
TITLE OF INVENTION: HUMAN SEF ISOFORMS AND METHODS OF USING SAME FOR CANCER GENE
FILE REFERENCE: 28385
CURRENT APPLICATION NUMBER: US/10/963,439
CURRENT FILING DATE: 2004-10-11
NUMBER OF SEQ ID NOS: 22
SOFTWARE: PatentIn version 3.3
SEQ ID NO 5
LENGTH: 739
TYPE: PRT
ORGANISM: Homo sapiens
US-10-963-439-5

Query Match 70.8%; Score 34; DB 9; Length 739;
Best Local Similarity 75.0%; Pred. No. 84;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 LPOLCTEL 9
Db 504 LPOLCSHL 511

RESULT 22

```

US-11-079-463-6485
; Sequence 6485, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FR
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6485
; LENGTH: 812
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-6485

Query Match          70.8%; Score 34; DB 11; Length 812;
Best Local Similarity 85.7%; Pred. No. 92;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 KLPOLCT 7
        |||||
Db      764 KLPOLCT 770

RESULT 23
US-11-188-298-4065
; Sequence 4065, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 4065
; LENGTH: 175
; TYPE: PRT
; ORGANISM: Trifolium aestivum
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(175)
; OTHER INFORMATION: unsure at all Xaa locations
US-11-188-298-4065

Query Match          68.8%; Score 33; DB 11; Length 175;
Best Local Similarity 85.7%; Pred. No. 35;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
        |||||
Db      65 LPOLCTE 71

RESULT 24
US-11-079-463-9899
; Sequence 9899, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FR
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; PRIOR FILING DATE: 2005-03-14

```

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; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 9899
; LENGTH: 243
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-9899

Query Match          68.8%; Score 33; DB 11; Length 243;
Best Local Similarity 71.4%; Pred. No. 47;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
        |||||
Db      144 VPOLCTE 150

RESULT 25
US-11-072-512-3188
; Sequence 3188, Application US/11072512
; Publication No. US20060029945A1
; GENERAL INFORMATION:
; APPLICANT: ISOGAI, TAKAO
; APPLICANT: SUGIYAMA, TOMOYASU
; APPLICANT: OTSUKI, TETSUJI
; APPLICANT: WAKAMATSU, AI
; APPLICANT: SATO, HIROYUKI
; APPLICANT: ISHII, SHIZUKO
; APPLICANT: YAMAMOTO, JUN-ICHI
; APPLICANT: ISONO, YUUKO
; APPLICANT: HIO, YURI
; APPLICANT: OTSUKA, KAORU
; APPLICANT: NAGAI, KEIICHI
; APPLICANT: IRIE, RYOTARO
; APPLICANT: TAMECHIKA, ICHIRO
; APPLICANT: SEKI, NAOHITO
; APPLICANT: YOSHIKAWA, TSUTOMU
; APPLICANT: OTSUKA, MOTOYUKI
; APPLICANT: NAGAHARI, KENJI
; APPLICANT: MASURO, YASUHIKO
; TITLE OF INVENTION: Novel full length cDNA
; FILE REFERENCE: 084335-0191
; CURRENT APPLICATION NUMBER: US/11/072,512
; PRIOR FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US 60/350,978
; PRIOR FILING DATE: 2002-01-25
; PRIOR APPLICATION NUMBER: JP 2001-379298
; PRIOR FILING DATE: 2001-11-05
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3188
; LENGTH: 384
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-072-512-3188

Query Match          68.8%; Score 33; DB 11; Length 384;
Best Local Similarity 75.0%; Pred. No. 72;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY      2 LPOLCTEL 9
        |||||
Db      348 LPOLCTSL 355

RESULT 26
US-11-045-004-2686
; Sequence 2686, Application US/11045004
; Publication No. US20060078901A1
; GENERAL INFORMATION:

```



```
APPLICANT: BUCHRIESSER, CARMEN
APPLICANT: FRANGEUL, LIONEL
APPLICANT: COUVE, ELISABETH
APPLICANT: RUSNIOT, CHRISTOPHE
APPLICANT: PSHT, HAFIDA
APPLICANT: DEHOUE, PIERRE
APPLICANT: DUSOURGET, OLIVIER
APPLICANT: CHETOUANI, FARID
APPLICANT: NEDJARI, HAFED
APPLICANT: GLASER, PHILIPPE
APPLICANT: KUNST, FRANCK
APPLICANT: COSSART, PASCAL
APPLICANT: DANIELS, JUSTIN
APPLICANT: GOEBEL, WERNER
APPLICANT: KREFT, JURGEN
APPLICANT: KUHN, MICHAEL
APPLICANT: NG, EVA
APPLICANT: VAZQUEZ-BOLAND, ANTONIO
APPLICANT: DOMINGUEZ-BERNAL, GUSTAVO
APPLICANT: GARRIDO-GARCIA, PATRICIA
APPLICANT: TERRERZ-MARTINEZ, ALBERTO
APPLICANT: AMEND, ALEXANDRA
APPLICANT: CHAKRABORTY, TRINAD
APPLICANT: DOMANN, EUGEN
APPLICANT: HAIN, THORSTEN
APPLICANT: BERCHER, PATRICK
APPLICANT: CHARBIT, ALAIN
APPLICANT: DURANT, LIONEL
APPLICANT: PEREZ-DIAZ, JOSE-CLAUDIO
APPLICANT: BAQUERO, FERNANDO
APPLICANT: GARCIA DEL PORTILLO, FRANCISCO
APPLICANT: GOMEZ-LOPEZ, NURIA
APPLICANT: MADUENIO, ENCARN
APPLICANT: PABLOS, BETRIZ DE
APPLICANT: WEHLAND, JURGEN
APPLICANT: KARST, UWE
APPLICANT: ENTIAN, KARL-DIETER
APPLICANT: HAUF, JORG
APPLICANT: ROSE, MATTHIAS
APPLICANT: VOSS, HAMUT
APPLICANT: TITLE OF INVENTION: LISTERIA MONOCYTOGENES GENOME, POLYPEPTIDES AND USES
APPLICANT: FILE REFERENCE: 05394.0018-02
APPLICANT: CURRENT APPLICATION NUMBER: US/11/045.004
APPLICANT: PRIOR FILING DATE: 2005-01-28
APPLICANT: PRIOR APPLICATION NUMBER: 10/637,657
APPLICANT: PRIOR FILING DATE: 2003-08-11
APPLICANT: PRIOR APPLICATION NUMBER: 10/257,023
APPLICANT: PRIOR FILING DATE: 2002-10-08
APPLICANT: PRIOR APPLICATION NUMBER: PCT/FR01/01118
APPLICANT: PRIOR FILING DATE: 2001-04-11
APPLICANT: PRIOR APPLICATION NUMBER: FR 00/04,629
APPLICANT: PRIOR FILING DATE: 2000-04-11
APPLICANT: NUMBER OF SEQ ID NOS: 2854
APPLICANT: SOFTWARE: PatentIn version 3.3
APPLICANT: SEQ ID NO 2686
APPLICANT: LENGTH: 215
APPLICANT: TYPE: PRT
APPLICANT: ORGANISM: Listeria monocytogenes
US-11-045-004-2686

Query Match 66.7%; Score 32; DB 11; Length 215;
Best Local Similarity 66.7%; Pred. No. 65;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

QY 1 KLPOLCTEL 9
DB 199 RLPOLCTEL 207

RESULT 27
US-11-124-368A-285
Sequence 285, Application US/11124368A
Publication No. US20050287559A1

```
GENERAL INFORMATION:
APPLICANT: Michele Cargill
APPLICANT: James J. Devlin
APPLICANT: May Luke
APPLICANT: TITLE OF INVENTION: Genetic Polymorphisms Associated with
APPLICANT: TITLE OF INVENTION: Vascular Diseases, Methods of Detection and Uses Thereof
APPLICANT: FILE REFERENCE: CL001524
APPLICANT: CURRENT APPLICATION NUMBER: US/11/124,368A
APPLICANT: CURRENT FILING DATE: 2005-05-09
APPLICANT: PRIOR APPLICATION NUMBER: US 60/568,845
APPLICANT: PRIOR FILING DATE: 2004-05-07
APPLICANT: PRIOR APPLICATION NUMBER: US 60/625,936
APPLICANT: PRIOR FILING DATE: 2004-11-09
APPLICANT: NUMBER OF SEQ ID NOS: 21112
APPLICANT: SOFTWARE: FastSeq for Windows Version 4.0
APPLICANT: SEQ ID NO 285
APPLICANT: LENGTH: 345
APPLICANT: TYPE: PRT
APPLICANT: ORGANISM: Homo sapiens
US-11-124-368A-285
```

Query Match 66.7%; Score 32; DB 11; Length 345;
Best Local Similarity 83.3%; Pred. No. 1e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 POLCTE 8
DB 320 POLCTE 325

```
RESULT 28
US-10-203-486-10
Sequence 10, Application US/10203486
Publication No. US20060035315A1
GENERAL INFORMATION:
APPLICANT: INCYTE GENOMICS, INC.
APPLICANT: YU, Henry
APPLICANT: TANG, Y. Tom
APPLICANT: LAL, Preeti
APPLICANT: POLICKY, Jennifer L.
APPLICANT: NGUYEN, Daniel B.
APPLICANT: NGUYEN, Janice
APPLICANT: YAO, Monique G.
APPLICANT: KHAN, Farrah A.
APPLICANT: WALIA, Ameena R.
APPLICANT: GANDHI, Arvind K.
APPLICANT: TRIBOULET, Catherine M.
APPLICANT: PATTERSON, Chandra
APPLICANT: THORNTON, Michael
APPLICANT: GREENE, Barrie D.
APPLICANT: HERNANDEZ, Roberto
APPLICANT: BOROWSKY, Mark L.
APPLICANT: APPLICANT: SANJAYWALA, Madhu S.
APPLICANT: TITLE OF INVENTION: TRANSPORTERS AND ION CHANNELS
APPLICANT: FILE REFERENCE: PI-0042 PCT
APPLICANT: CURRENT APPLICATION NUMBER: US/10/203,486
APPLICANT: CURRENT FILING DATE: 2002-08-08
APPLICANT: NUMBER OF SEQ ID NOS: 26
APPLICANT: SOFTWARE: PERL Program
APPLICANT: SEQ ID NO 10
APPLICANT: LENGTH: 473
APPLICANT: TYPE: PRT
APPLICANT: ORGANISM: Homo sapiens
APPLICANT: FEATURE:
APPLICANT: NAME/KEY: misc feature
APPLICANT: OTHER INFORMATION: Incyte ID No: 4297813CD1
US-10-203-486-10

Query Match 66.7%; Score 32; DB 9; Length 473;
Best Local Similarity 75.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

QY 2 LPOLCTEL 9

Db 398 LPQLCPSEL 405

RESULT 29

US-10-467-7682
; Sequence 7682, Application US/10467657
; Publication No. US20050260581A1
; GENERAL INFORMATION:
; APPLICANT: CHIRON SPA
; APPLICANT: FONTANA Maria Rita
; APPLICANT: PIZZA Mariagrazia
; APPLICANT: MASTIGNANI Vega
; APPLICANT: MONACI Elisabetta
; TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
; FILE REFERENCE:
; CURRENT APPLICATION NUMBER: US/10/467,657
; PRIOR FILING DATE: 2003-08-11
; PRIOR APPLICATION NUMBER: GB-0103424.8
; PRIOR FILING DATE: 2001-02-12
; NUMBER OF SEQ ID NOS: 9218
; SOFTWARE: SeqWin39, version 1.04
; SEQ ID NO 7682
; LENGTH: 501
; TYPE: PRT
; ORGANISM: Neisseria gonorrhoeae
US-10-467-657-7682

Query Match 66.7%; Score 32; DB 9; Length 501;
Best Local Similarity 55.6%; Pred. No. 1.4e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
:|||||
Db 129 QLPKICREL 137

RESULT 30
US-11-103-957-61
; Sequence 61, Application US/11103957
; Publication No. US20050281847A1
; GENERAL INFORMATION:
; APPLICANT: Berthet, Francois-Xavier Jacques
; APPLICANT: Lobet, Yves
; APPLICANT: Poolman, Jan
; APPLICANT: Verlant, Vincent Georges Christian Louis
; TITLE OF INVENTION: Vaccine Composition
; FILE REFERENCE: B45261
; CURRENT APPLICATION NUMBER: US/11/103,957
; CURRENT FILING DATE: 2005-04-12
; PRIOR APPLICATION NUMBER: US/10/467,534
; PRIOR FILING DATE: 2004-02-03
; PRIOR APPLICATION NUMBER: PCT/EP02/01356
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: GB 0103169.9
; PRIOR FILING DATE: 2001-02-08
; NUMBER OF SEQ ID NOS: 108
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 61
; LENGTH: 553
; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-11-103-957-61

Query Match 66.7%; Score 32; DB 11; Length 553;
Best Local Similarity 66.7%; Pred. No. 1.6e+02;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
:|||||
Db 12 KTPSLCREL 20

RESULT 31

US-10-203-486-8
; Sequence 8, Application US/10203486
; Publication No. US20060035315A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE GENOMICS, INC.
; APPLICANT: YUE, Henry
; APPLICANT: TANG, Y. Tom
; APPLICANT: LAL, Preeti
; APPLICANT: POLICKY, Jennifer L.
; APPLICANT: NGUYEN, Daniel B.
; APPLICANT: AU-YOUNG, Janice
; APPLICANT: YAO, Montique G.
; APPLICANT: KHAN, Farhan A.
; APPLICANT: WALIA, Narender K.
; APPLICANT: GANDHI, Ameena R.
; APPLICANT: TRIBOULET, Catherine M.
; APPLICANT: PATTERSON, Chandra
; APPLICANT: THORNTON, Michael
; APPLICANT: GREENE, Barrie D.
; APPLICANT: HERNANDEZ, Roberto
; APPLICANT: BOROWSKY, Mark L.
; APPLICANT: SANTANALIA, Madhu S.
; TITLE OF INVENTION: TRANSPORTERS AND ION CHANNELS
; FILE REFERENCE: PI-0042 PCT
; CURRENT APPLICATION NUMBER: US/10/203,486
; CURRENT FILING DATE: 2002-08-08
; NUMBER OF SEQ ID NOS: 26
; SOFTWARE: PERL Program
; SEQ ID NO 8
; LENGTH: 577
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No: 7472477CD1
US-10-203-486-8

Query Match 66.7%; Score 32; DB 9; Length 577;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LPOLCTEL 9
:|||||
Db 502 LPQLCPSEL 509

RESULT 32
US-11-098-686-10124
; Sequence 10124, Application US/11098686
; Publication No. US20060024696A1
; GENERAL INFORMATION:
; APPLICANT: Kapur, Vivek and Gebhart, Connie J.
; TITLE OF INVENTION: NUCLEIC ACID AND POLYPEPTIDE SEQUENCES
; TITLE OF INVENTION: FROM LAWSONIA INTRACELLULARIS AND METHODS OF USING
; FILE REFERENCE: 09531-128001
; CURRENT APPLICATION NUMBER: US/11/098,686
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31318
; PRIOR FILING DATE: 2003-10-01
; PRIOR APPLICATION NUMBER: US 60/416,395
; PRIOR FILING DATE: 2002-10-04
; NUMBER OF SEQ ID NOS: 11433
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10124
; LENGTH: 626
; TYPE: PRT
; ORGANISM: Lawsonia intracellularis
US-11-098-686-10124

Query Match 66.7%; Score 32; DB 11; Length 626;
Best Local Similarity 55.6%; Pred. No. 1.8e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KLPOCTEL 9
Db 167 RYPOFCKL 175

RESULT 33
US-11-079-463-6039
; Sequence 6039, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FR
; FILE REFERENCE: PAT00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6039
; LENGTH: 686
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-6039

Query Match 66.7%; Score 31; DB 11; Length 686;
Best Local Similarity 55.6%; Pred. No. 1.9e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 KLPOCTEL 9
Db 64 QLEVCHEL 72

RESULT 34
US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/1004137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 26
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 68
US-10-530-253-26

Query Match 64.6%; Score 31; DB 9; Length 158;
Best Local Similarity 66.7%; Pred. No. 76;
Matches 6; Conservative 0; Mismatches 3; Indels 0; Gaps 0;

Qy 1 KLPOCTEL 9
Db 13 KLPOCTEL 21

RESULT 35
US-11-079-463-7727

; Sequence 7727, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FR
; FILE REFERENCE: PAT00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 7727
; LENGTH: 197
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-7727

Query Match 64.6%; Score 31; DB 11; Length 197;
Best Local Similarity 55.6%; Pred. No. 93;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KLPOCTEL 9
Db 138 KLPERCREI 146

RESULT 36
US-11-096-568A-26047
; Sequence 26047, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 26047
; LENGTH: 206
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)-(206)
; OTHER INFORMATION: Ceres Seq. ID no. 13498304
US-11-096-568A-26047

Query Match 64.6%; Score 31; DB 11; Length 206;
Best Local Similarity 66.7%; Pred. No. 97;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 KLPOCTEL 9
Db 88 KLSQLCDDL 96

RESULT 37
US-11-087-099-2315
; Sequence 2315, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 2315
; LENGTH: 209

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; TYPE: PRT
; ORGANISM: Zea mays
US-11-087-099-2315

Query Match
Best Local Similarity 64.6%; Score 31; DB 11; Length 209;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 KLPOLCT 7
Db 20 KLPOLCT 26

RESULT 38
US-11-096-568A-26045
; Sequence 26045, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 26045
; LENGTH: 226
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(226)
; OTHER INFORMATION: Ceres Seq. ID no. 13498302
US-11-096-568A-26045

Query Match
Best Local Similarity 64.6%; Score 31; DB 11; Length 226;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 KLPOLCTEL 9
Db 108 KLSQICDDL 116

RESULT 39
US-10-242-586-8
; Sequence 8, Application US/10242586
; Publication No. US20060073548A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Eaton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Philippe
; APPLICANT: Watanabe, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C21
; CURRENT APPLICATION NUMBER: US/10/242,586
; CURRENT FILING DATE: 2002-09-11
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
```

```
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-242-586-8

Query Match
Best Local Similarity 64.6%; Score 31; DB 9; Length 229;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 LPOLCTE 8
Db 31 LPOLCTQ 37

RESULT 40
US-10-242-902-8
; Sequence 8, Application US/10242902
; Publication No. US20060073549A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Eaton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Philippe
; APPLICANT: Watanabe, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C54
; CURRENT APPLICATION NUMBER: US/10/242,902
; CURRENT FILING DATE: 2002-09-13
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
```

```
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-902-8
```

```
Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 LPOLCTE 8
        ||::||:
Db       31 LPEICTQ 37
```

```
RESULT 41
US-10-243-116-8
; Sequence 8, Application US/10243116
; Publication No. US20060073550A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Philippe
; APPLICANT: Watande, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C38
; CURRENT FILING DATE: 2002-09-13
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-116-8
```

```
Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 LPOLCTE 8
        ||::||:
Db       31 LPEICTQ 37
```

```
RESULT 42
US-10-243-136-8
; Sequence 8, Application US/10243136
; Publication No. US20060074228A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Philippe
; APPLICANT: Watande, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
```

```
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C32
; CURRENT FILING DATE: 2002-09-12
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-136-8
```

```
Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 LPOLCTE 8
        ||::||:
Db       31 LPEICTQ 37
```

```
RESULT 43
US-10-243-189-8
; Sequence 8, Application US/10243189
; Publication No. US20060074033A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
```

```
/ APPLICANT: Stephan,Jean-Philippe
/ APPLICANT: Watanabe,Colin
/ APPLICANT: Wood,William
/ APPLICANT: Zhang,Zemin
/ APPLICANT: Fong,Sherman
/ TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
/ TITLE OF INVENTION: ACIDS ENCODING THE SAME
/ FILE REFERENCE: P3630R1C44
/ CURRENT APPLICATION NUMBER: US/10/243,189
/ PRIOR FILING DATE: 2002-09-12
/ PRIOR APPLICATION NUMBER: 10/197942
/ PRIOR FILING DATE: 2002-07-18
/ PRIOR APPLICATION NUMBER: 60/059114
/ PRIOR FILING DATE: 1997-09-17
/ PRIOR APPLICATION NUMBER: 60/063046
/ PRIOR FILING DATE: 1997-10-24
/ PRIOR APPLICATION NUMBER: 60/065027
/ PRIOR FILING DATE: 1997-11-10
/ PRIOR APPLICATION NUMBER: 60/079689
/ PRIOR FILING DATE: 1998-03-27
/ PRIOR APPLICATION NUMBER: 60/086478
/ PRIOR FILING DATE: 1998-05-22
/ PRIOR APPLICATION NUMBER: 60/087607
/ PRIOR FILING DATE: 1998-06-02
/ PRIOR APPLICATION NUMBER: 60/089801
/ PRIOR FILING DATE: 1998-06-18
/ PRIOR APPLICATION NUMBER: 60/090557
/ PRIOR FILING DATE: 1998-06-24
/ PRIOR APPLICATION NUMBER: 60/090689
/ PRIOR FILING DATE: 1998-06-25
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 116
/ SEQ ID NO 8
/ LENGTH: 229
/ TYPE: PRT
/ ORGANISM: Homo Sapien
US-10-243-189-8

Query Match          64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
DB      31 LPEICTQ 37

RESULT 44
US-10-243-215-8
/ Sequence 8, Application US/10243215
/ Publication No. US20060073551A1
/ GENERAL INFORMATION:
/ APPLICANT: Baker, Kevin
/ APPLICANT: Baton, Dan
/ APPLICANT: Filvaroff, Ellen
/ APPLICANT: Goddard, Audrey
/ APPLICANT: Grimaldi, J. Christopher
/ APPLICANT: Gurney, Austin
/ APPLICANT: Smith, Victoria
/ APPLICANT: Stephan, Jean-Philippe
/ APPLICANT: Watanabe, Colin
/ APPLICANT: Wood, William
/ APPLICANT: Zhang, Zemin
/ APPLICANT: Fong, Sherman
/ TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
/ TITLE OF INVENTION: ACIDS ENCODING THE SAME
/ FILE REFERENCE: P3630R1C27
/ CURRENT APPLICATION NUMBER: US/10/243,215
/ PRIOR FILING DATE: 2002-09-12
/ PRIOR APPLICATION NUMBER: 10/197942
/ PRIOR FILING DATE: 2002-07-18
/ PRIOR APPLICATION NUMBER: 60/059114
/ PRIOR FILING DATE: 1997-09-17
```

```
/ PRIOR APPLICATION NUMBER: 60/063046
/ PRIOR FILING DATE: 1997-10-24
/ PRIOR APPLICATION NUMBER: 60/065027
/ PRIOR FILING DATE: 1997-11-10
/ PRIOR APPLICATION NUMBER: 60/079689
/ PRIOR FILING DATE: 1998-03-27
/ PRIOR APPLICATION NUMBER: 60/086478
/ PRIOR FILING DATE: 1998-05-22
/ PRIOR APPLICATION NUMBER: 60/087607
/ PRIOR FILING DATE: 1998-06-02
/ PRIOR APPLICATION NUMBER: 60/089801
/ PRIOR FILING DATE: 1998-06-18
/ PRIOR APPLICATION NUMBER: 60/090689
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 116
/ SEQ ID NO 8
/ LENGTH: 229
/ TYPE: PRT
/ ORGANISM: Homo Sapien
US-10-243-215-8

Query Match          64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
DB      31 LPEICTQ 37

RESULT 45
US-10-243-236-8
/ Sequence 8, Application US/10243236
/ Publication No. US20060073552A1
/ GENERAL INFORMATION:
/ APPLICANT: Baker, Kevin
/ APPLICANT: Baton, Dan
/ APPLICANT: Filvaroff, Ellen
/ APPLICANT: Goddard, Audrey
/ APPLICANT: Grimaldi, J. Christopher
/ APPLICANT: Gurney, Austin
/ APPLICANT: Smith, Victoria
/ APPLICANT: Stephan, Jean-Philippe
/ APPLICANT: Watanabe, Colin
/ APPLICANT: Wood, William
/ APPLICANT: Zhang, Zemin
/ APPLICANT: Fong, Sherman
/ TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
/ TITLE OF INVENTION: ACIDS ENCODING THE SAME
/ FILE REFERENCE: P3630R1C35
/ CURRENT APPLICATION NUMBER: US/10/243,236
/ PRIOR FILING DATE: 2002-09-12
/ PRIOR APPLICATION NUMBER: 10/197942
/ PRIOR FILING DATE: 2002-07-18
/ PRIOR APPLICATION NUMBER: 60/059114
/ PRIOR FILING DATE: 1997-09-17
/ PRIOR APPLICATION NUMBER: 60/063046
/ PRIOR FILING DATE: 1997-10-24
/ PRIOR APPLICATION NUMBER: 60/065027
/ PRIOR FILING DATE: 1997-11-10
/ PRIOR APPLICATION NUMBER: 60/079689
/ PRIOR FILING DATE: 1998-03-27
/ PRIOR APPLICATION NUMBER: 60/086478
/ PRIOR FILING DATE: 1998-05-22
/ PRIOR APPLICATION NUMBER: 60/087607
/ PRIOR FILING DATE: 1998-06-02
/ PRIOR APPLICATION NUMBER: 60/089801
/ PRIOR FILING DATE: 1998-06-18
/ PRIOR APPLICATION NUMBER: 60/090557
/ PRIOR FILING DATE: 1998-06-24
```

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; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-236-8

Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
Db      31 LPEICTQ 37

RESULT 46
US-10-243-298-8
; Sequence 8, Application US/10243298
; Publication No. US20060073553A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Phillippe
; APPLICANT: Watanabe, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C49
; CURRENT APPLICATION NUMBER: US/10/243,298
; PRIOR FILING DATE: 2002-09-13
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-298-8

Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```

QY      2 LPOLCTE 8
Db      31 LPEICTQ 37

RESULT 47
US-10-243-304-8
; Sequence 8, Application US/10243304
; Publication No. US20060073554A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
; APPLICANT: Grimaldi, J. Christopher
; APPLICANT: Gurney, Austin
; APPLICANT: Smith, Victoria
; APPLICANT: Stephan, Jean-Phillippe
; APPLICANT: Watanabe, Colin
; APPLICANT: Wood, William
; APPLICANT: Zhang, Zemin
; APPLICANT: Fong, Sherman
; TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
; FILE REFERENCE: P3630R1C42
; CURRENT APPLICATION NUMBER: US/10/243,304
; PRIOR FILING DATE: 2002-09-12
; PRIOR APPLICATION NUMBER: 10/197942
; PRIOR FILING DATE: 2002-07-18
; PRIOR APPLICATION NUMBER: 60/059114
; PRIOR FILING DATE: 1997-09-17
; PRIOR APPLICATION NUMBER: 60/063046
; PRIOR FILING DATE: 1997-10-24
; PRIOR APPLICATION NUMBER: 60/065027
; PRIOR FILING DATE: 1997-11-10
; PRIOR APPLICATION NUMBER: 60/079689
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: 60/086478
; PRIOR FILING DATE: 1998-05-22
; PRIOR APPLICATION NUMBER: 60/087607
; PRIOR FILING DATE: 1998-06-02
; PRIOR APPLICATION NUMBER: 60/089801
; PRIOR FILING DATE: 1998-06-18
; PRIOR APPLICATION NUMBER: 60/090557
; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-304-8

Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY      2 LPOLCTE 8
Db      31 LPEICTQ 37

RESULT 48
US-10-243-338-8
; Sequence 8, Application US/10243338
; Publication No. US20060073579A1
; GENERAL INFORMATION:
; APPLICANT: Baker, Kevin
; APPLICANT: Baton, Dan
; APPLICANT: Filvaroff, Ellen
; APPLICANT: Goddard, Audrey
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APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Philippe
APPLICANT: Watande, Colin
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
APPLICANT: Fong, Sherman
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3630R1C52
CURRENT APPLICATION NUMBER: US/10/243,338
PRIORITY FILING DATE: 2002-09-13
PRIORITY FILING DATE: 10/197942
PRIORITY FILING DATE: 2002-07-18
PRIORITY FILING DATE: 60/059114
PRIORITY FILING DATE: 1997-09-17
PRIORITY FILING DATE: 60/063046
PRIORITY FILING DATE: 1997-10-24
PRIORITY FILING DATE: 60/065027
PRIORITY FILING DATE: 1997-11-10
PRIORITY FILING DATE: 60/079689
PRIORITY FILING DATE: 1998-03-27
PRIORITY FILING DATE: 60/086478
PRIORITY FILING DATE: 1998-05-22
PRIORITY FILING DATE: 60/087607
PRIORITY FILING DATE: 1998-06-02
PRIORITY FILING DATE: 60/089801
PRIORITY FILING DATE: 1998-06-18
PRIORITY FILING DATE: 60/090557
PRIORITY FILING DATE: 1998-06-24
PRIORITY FILING DATE: 60/090689
PRIORITY FILING DATE: 1998-06-25
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 116
SEQ ID NO 8
LENGTH: 229
TYPE: PRT
ORGANISM: Homo Sapien
US-10-243-338-8
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Query Match 64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1,1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
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QY 2 LPOLCITE 8
DB 31 LPEICTQ 37
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RESULT 49
US-10-243-345-8
Sequence 8, Application US/10243345
Publication No. US20060073555A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Baton, Dan
APPLICANT: Filvaroff, Ellen
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Philippe
APPLICANT: Watande, Colin
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
APPLICANT: Fong, Sherman
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3630R1C3
CURRENT APPLICATION NUMBER: US/10/243,345
PRIORITY FILING DATE: 2002-09-12
PRIORITY FILING DATE: 10/197942
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PRIORITY FILING DATE: 2002-07-18
PRIORITY APPLICATION NUMBER: 60/059114
PRIORITY FILING DATE: 1997-09-17
PRIORITY APPLICATION NUMBER: 60/063046
PRIORITY FILING DATE: 1997-10-24
PRIORITY APPLICATION NUMBER: 60/065027
PRIORITY FILING DATE: 1997-11-10
PRIORITY APPLICATION NUMBER: 60/079689
PRIORITY FILING DATE: 1998-03-27
PRIORITY APPLICATION NUMBER: 60/086478
PRIORITY FILING DATE: 1998-05-22
PRIORITY APPLICATION NUMBER: 60/087607
PRIORITY FILING DATE: 1998-06-02
PRIORITY APPLICATION NUMBER: 60/089801
PRIORITY FILING DATE: 1998-06-18
PRIORITY APPLICATION NUMBER: 60/090557
PRIORITY FILING DATE: 1998-06-24
PRIORITY APPLICATION NUMBER: 60/090689
PRIORITY FILING DATE: 1998-06-25
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 116
SEQ ID NO 8
LENGTH: 229
TYPE: PRT
ORGANISM: Homo Sapien
US-10-243-345-8
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QY 2 LPOLCITE 8
DB 31 LPEICTQ 37
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RESULT 50
US-10-243-357-8
Sequence 8, Application US/10243357
Publication No. US20060073555A1
GENERAL INFORMATION:
APPLICANT: Baker, Kevin
APPLICANT: Baton, Dan
APPLICANT: Filvaroff, Ellen
APPLICANT: Goddard, Audrey
APPLICANT: Grimaldi, J. Christopher
APPLICANT: Gurney, Austin
APPLICANT: Smith, Victoria
APPLICANT: Stephan, Jean-Philippe
APPLICANT: Watande, Colin
APPLICANT: Wood, William
APPLICANT: Zhang, Zemin
APPLICANT: Fong, Sherman
TITLE OF INVENTION: SECRETED AND TRANSMEMBRANE POLYPEPTIDES AND NUCLEIC
FILE REFERENCE: P3630R1C55
CURRENT APPLICATION NUMBER: US/10/243,357
PRIORITY FILING DATE: 2002-09-13
PRIORITY APPLICATION NUMBER: 10/197942
PRIORITY FILING DATE: 2002-07-18
PRIORITY APPLICATION NUMBER: 60/059114
PRIORITY FILING DATE: 1997-09-17
PRIORITY APPLICATION NUMBER: 60/063046
PRIORITY FILING DATE: 1997-10-24
PRIORITY APPLICATION NUMBER: 60/065027
PRIORITY FILING DATE: 1997-11-10
PRIORITY APPLICATION NUMBER: 60/079689
PRIORITY FILING DATE: 1998-03-27
PRIORITY APPLICATION NUMBER: 60/086478
PRIORITY FILING DATE: 1998-05-22
PRIORITY APPLICATION NUMBER: 60/087607
PRIORITY FILING DATE: 1998-06-02
PRIORITY APPLICATION NUMBER: 60/089801
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; PRIOR FILING DATE: 1998-06-18
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; PRIOR FILING DATE: 1998-06-24
; PRIOR APPLICATION NUMBER: 60/090689
; PRIOR FILING DATE: 1998-06-25
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 116
; SEQ ID NO 8
; LENGTH: 229
; TYPE: PRT
; ORGANISM: Homo Sapien
US-10-243-357-8
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Query Match      64.6%; Score 31; DB 9; Length 229;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
QY      2 LPQLCTE 8
      ||::||:
Db      31 LPEICTQ 37
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Job time : 9.4 secs
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GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 05:36:41 ; Search time 20.9 Seconds
(without alignments)
35.602 Million cell updates/sec

Title: US-08-170-344-3
Perfect score: 47
Sequence: 1 QLCLEQTT 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Issued Patents, AA:
1: /cgn2_6/ptodata/1/1aa/5-COMB.pep:*
2: /cgn2_6/ptodata/1/1aa/6-COMB.pep:*
3: /cgn2_6/ptodata/1/1aa/H-COMB.pep:*
4: /cgn2_6/ptodata/1/1aa/BCTUS-COMB.pep:*
5: /cgn2_6/ptodata/1/1aa/BB-COMB.pep:*
6: /cgn2_6/ptodata/1/1aa/Backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	47	100.0	30	1	US-08-363-586-4
3	47	100.0	30	2	US-09-980-523A-4
4	47	100.0	151	2	US-09-701-080C-18
5	47	100.0	158	2	US-09-980-523A-2
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7	47	100.0	162	1	US-08-316-239B-4
8	47	100.0	172	2	US-08-860-165-14
9	47	100.0	172	2	US-09-359-382-14
10	47	100.0	182	1	US-08-117-083-10
11	47	100.0	243	2	US-09-462-993-1
12	47	100.0	266	2	US-08-860-165-10
13	47	100.0	266	2	US-09-359-382-10
14	47	100.0	266	2	US-09-367-309A-1
15	47	100.0	273	2	US-09-485-885-4
16	47	100.0	292	2	US-09-485-885-10
17	47	100.0	371	2	US-09-485-885-6
18	47	100.0	390	2	US-09-485-885-14
19	34	72.3	4019	2	US-09-854-133-425
20	33	70.2	32	1	US-08-466-283-2
21	33	70.2	32	2	US-08-164-768-2
22	33	70.2	141	2	US-09-270-767-33527
23	33	70.2	141	2	US-09-270-767-48744
24	33	70.2	158	1	US-08-347-904B-19
25	33	70.2	158	2	US-08-767-942A-19
26	33	70.2	271	1	US-08-117-083-14
27	33	70.2	278	2	US-09-485-885-21

28	33	70.2	383	2	US-09-485-885-23	Sequence 23, Appl
29	33	70.2	1098	2	US-10-104-047-2475	Sequence 2475, Ap
30	33	70.2	1104	2	US-09-793-998-11	Sequence 11, Appl
31	32	68.1	9	1	US-08-787-547-101	Sequence 101, App
32	32	68.1	9	2	US-09-601-729-274	Sequence 274, App
33	32	68.1	203	2	US-09-248-796A-17743	Sequence 17743, A
34	32	68.1	211	2	US-09-543-681A-8111	Sequence 8111, Ap
35	32	68.1	236	2	US-09-634-137-32	Sequence 32, Appl
36	32	68.1	251	2	US-09-270-767-33015	Sequence 33015, A
37	32	68.1	469	2	US-09-543-681A-5423	Sequence 5423, Ap
38	32	68.1	771	2	US-09-121-964-9	Sequence 9, Appl
39	31	66.0	9	2	US-08-159-339A-320	Sequence 320, App
40	31	66.0	14	2	US-08-271-539-3	Sequence 3, Appl
41	31	66.0	56	2	US-09-513-999C-6885	Sequence 6885, Ap
42	31	66.0	108	1	US-08-234-812-3	Sequence 3, Appl
43	31	66.0	108	1	US-08-663-809-3	Sequence 3, Appl
44	31	66.0	129	1	US-08-049-503-1	Sequence 1, Appl
45	31	66.0	129	1	US-08-225-224-2	Sequence 2, Appl
46	31	66.0	129	1	US-08-470-299-21	Sequence 21, Appl
47	31	66.0	129	1	US-08-874-697-1	Sequence 1, Appl
48	31	66.0	129	2	US-08-722-258-2	Sequence 2, Appl
49	31	66.0	129	2	US-08-897-020-1	Sequence 1, Appl
50	31	66.0	129	2	US-08-765-012A-16	Sequence 16, Appl
51	31	66.0	129	2	US-08-765-012A-17	Sequence 17, Appl
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54	31	66.0	129	2	US-08-765-012A-20	Sequence 20, Appl
55	31	66.0	129	2	US-09-350-823-1	Sequence 1, Appl
56	31	66.0	129	2	US-09-408-629-1	Sequence 1, Appl
57	31	66.0	129	2	US-08-271-539-46	Sequence 46, Appl
58	31	66.0	129	2	US-09-462-941-11	Sequence 11, Appl
59	31	66.0	129	2	US-09-609-027B-3	Sequence 3, Appl
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61	31	66.0	130	2	US-08-765-012A-11	Sequence 11, Appl
62	31	66.0	130	2	US-08-765-012A-12	Sequence 12, Appl
63	31	66.0	130	2	US-08-765-012A-13	Sequence 13, Appl
64	31	66.0	130	2	US-08-765-012A-14	Sequence 14, Appl
65	31	66.0	130	2	US-08-765-012A-15	Sequence 15, Appl
66	31	66.0	130	2	US-08-765-012A-16	Sequence 16, Appl
67	31	66.0	131	2	US-08-765-012A-17	Sequence 17, Appl
68	31	66.0	131	2	US-08-765-012A-18	Sequence 18, Appl
69	31	66.0	131	2	US-08-765-012A-19	Sequence 19, Appl
70	31	66.0	131	2	US-08-765-012A-20	Sequence 20, Appl
71	31	66.0	133	1	US-08-765-012A-10	Sequence 10, Appl
72	31	66.0	133	1	US-08-224-812-2	Sequence 2, Appl
73	31	66.0	133	1	US-08-663-809-2	Sequence 2, Appl
74	31	66.0	133	2	US-08-759-628-7	Sequence 7, Appl
75	31	66.0	153	2	US-09-522-217-112	Sequence 112, App
76	31	66.0	153	2	US-09-923-246-112	Sequence 112, App
77	31	66.0	153	2	US-10-295-723-112	Sequence 112, App
78	31	66.0	153	2	US-10-282-622-8	Sequence 8, Appl
79	31	66.0	165	2	US-09-949-016-7928	Sequence 7928, Ap
80	31	66.0	214	1	US-08-766-605-1	Sequence 1, Appl
81	31	66.0	214	1	US-09-094-212-1	Sequence 1, Appl
82	31	66.0	226	2	US-09-489-039A-7406	Sequence 7406, Ap
83	31	66.0	300	2	US-08-897-340-32	Sequence 32, Appl
84	31	66.0	300	2	US-09-252-329-32	Sequence 32, Appl
85	31	66.0	346	1	US-09-543-681A-5716	Sequence 6716, Ap
86	31	66.0	382	1	US-08-470-299-7	Sequence 7, Appl
87	31	66.0	382	1	US-08-470-299-10	Sequence 10, Appl
88	31	66.0	387	1	US-08-470-299-4	Sequence 4, Appl
89	31	66.0	498	2	US-09-232-468A-18	Sequence 18, Appl
90	31	66.0	498	2	US-08-686-968C-231	Sequence 231, App
91	31	66.0	498	2	US-09-784-984B-32	Sequence 52, Appl
92	31	66.0	637	2	US-09-187-906-7	Sequence 7, Appl
93	31	66.0	826	2	US-09-830-762-5	Sequence 5, Appl
94	31	66.0	826	2	US-08-705-660-34	Sequence 34, Appl
95	31	66.0	826	2	US-08-989-043-34	Sequence 34, Appl
96	31	66.0	3224	1	US-09-538-092-1161	Sequence 1161, Ap
97	31	66.0	3224	2	US-09-315-355A-34	Sequence 34, Appl
98	31	66.0	3224	2	US-08-159-339A-129	Sequence 329, App
99	30	63.8	10	2	US-08-159-339A-331	Sequence 331, App
100	30	63.8	10	2	US-08-159-339A-331	Sequence 331, App

101	30	63.8	31	1	US-08-438-753B-7	Sequence 7, Appl1	174	29	61.7	14	1	US-07-909-122-3	Sequence 3, Appl1
102	30	63.8	31	1	US-08-443-883A-7	Sequence 7, Appl1	175	29	61.7	50	2	US-09-621-976-4272	Sequence 4772, Ap
103	30	63.8	31	1	US-08-631-328-7	Sequence 7, Appl1	176	29	61.7	50	2	US-09-621-976-7318	Sequence 7318, Ap
104	30	63.8	31	1	US-08-453-524B-7	Sequence 7, Appl1	177	29	61.7	120	2	US-10-101-464A-679	Sequence 679, Ap
105	30	63.8	31	1	US-08-045-021B-7	Sequence 7, Appl1	178	29	61.7	137	2	US-09-328-352-6431	Sequence 6431, Ap
106	30	63.8	31	2	US-09-045-467-7	Sequence 7, Appl1	179	29	61.7	165	1	US-08-462-894-4	Sequence 4, Appl1
107	30	63.8	56	2	US-08-866-545-10	Sequence 10, Appl1	180	29	61.7	165	1	US-08-206-185-4	Sequence 4, Appl1
108	30	63.8	58	2	US-09-627-775-10	Sequence 10, Appl1	181	29	61.7	166	1	US-08-462-894-2	Sequence 2, Appl1
109	30	63.8	62	2	US-09-902-540-16179	Sequence 16179, A	182	29	61.7	166	1	US-08-462-894-6	Sequence 6, Appl1
110	30	63.8	90	2	US-09-107-433-4444	Sequence 4444, Ap	183	29	61.7	166	1	US-08-206-185-2	Sequence 2, Appl1
111	30	63.8	94	2	US-09-270-767-59551	Sequence 59551, A	184	29	61.7	166	1	US-08-206-185-6	Sequence 6, Appl1
112	30	63.8	101	2	US-09-270-767-44130	Sequence 44130, A	185	29	61.7	170	1	US-08-462-894-12	Sequence 12, Appl1
113	30	63.8	120	2	US-09-489-039A-12045	Sequence 12045, A	186	29	61.7	170	1	US-08-206-185-11	Sequence 11, Appl1
114	30	63.8	135	2	US-09-902-540-14438	Sequence 14438, A	187	29	61.7	171	1	US-08-462-894-11	Sequence 11, Appl1
115	30	63.8	138	2	US-08-828-683A-20	Sequence 20, Appl1	188	29	61.7	171	1	US-08-462-894-13	Sequence 13, Appl1
116	30	63.8	141	2	US-08-906-769-135	Sequence 135, App	189	29	61.7	171	1	US-08-206-185-11	Sequence 11, Appl1
117	30	63.8	141	2	US-08-906-616-135	Sequence 135, App	190	29	61.7	171	1	US-08-206-185-13	Sequence 13, Appl1
118	30	63.8	141	2	US-08-639-075A-135	Sequence 135, App	191	29	61.7	171	1	US-08-793-294-2	Sequence 2, Appl1
119	30	63.8	141	2	US-09-012-431-135	Sequence 135, App	192	29	61.7	172	1	US-08-793-294-1	Sequence 1, Appl1
120	30	63.8	141	2	US-09-012-682-135	Sequence 135, App	193	29	61.7	172	1	US-08-793-294-3	Sequence 3, Appl1
121	30	63.8	141	2	US-08-906-613-135	Sequence 135, App	194	29	61.7	172	1	US-08-462-894-10	Sequence 10, Appl1
122	30	63.8	141	2	US-08-438-753B-2	Sequence 2, Appl1	195	29	61.7	188	1	US-08-206-185-10	Sequence 10, Appl1
123	30	63.8	172	1	US-08-443-883A-2	Sequence 2, Appl1	196	29	61.7	189	1	US-08-462-894-8	Sequence 8, Appl1
124	30	63.8	172	1	US-08-631-328-2	Sequence 2, Appl1	197	29	61.7	189	1	US-08-462-894-8	Sequence 8, Appl1
125	30	63.8	172	1	US-08-455-524B-2	Sequence 2, Appl1	198	29	61.7	189	1	US-08-206-185-8	Sequence 8, Appl1
126	30	63.8	172	1	US-08-455-021B-2	Sequence 2, Appl1	199	29	61.7	199	1	US-09-583-110-4167	Sequence 4167, Ap
127	30	63.8	172	2	US-09-045-467-2	Sequence 2, Appl1	200	29	61.7	199	1	US-09-107-433-4096	Sequence 4096, Ap
128	30	63.8	172	2	US-08-934-395A-18	Sequence 18, Appl1	201	29	61.7	273	2	US-09-328-352-7386	Sequence 7386, Ap
129	30	63.8	172	2	US-08-616-904-2	Sequence 2, Appl1	202	29	61.7	349	2	US-09-949-016-7726	Sequence 7726, Ap
130	30	63.8	172	2	US-09-599-413-2	Sequence 2, Appl1	203	29	61.7	444	2	US-10-363-937-14	Sequence 14, Appl1
131	30	63.8	172	2	US-09-599-413-4	Sequence 4, Appl1	204	29	61.7	440	2	US-09-461-325-250	Sequence 250, App
132	30	63.8	172	2	US-09-599-413-5	Sequence 5, Appl1	205	29	61.7	490	2	US-10-012-542-250	Sequence 250, App
133	30	63.8	172	2	US-09-599-413-6	Sequence 6, Appl1	206	29	61.7	490	2	US-10-012-542-518	Sequence 518, App
134	30	63.8	172	2	US-09-599-413-7	Sequence 7, Appl1	207	29	61.7	490	2	US-10-115-123-250	Sequence 250, App
135	30	63.8	172	2	US-09-599-413-8	Sequence 8, Appl1	208	29	61.7	490	2	US-10-115-123-518	Sequence 518, App
136	30	63.8	172	2	US-09-599-413-9	Sequence 9, Appl1	209	29	61.7	492	2	US-09-794-236-2	Sequence 2, Appl1
137	30	63.8	172	2	US-09-599-413-10	Sequence 10, Appl1	210	29	61.7	492	2	US-09-345-469-1	Sequence 1, Appl1
138	30	63.8	172	2	US-09-599-413-18	Sequence 18, Appl1	211	29	61.7	515	2	US-09-461-325-219	Sequence 219, App
139	30	63.8	172	2	US-09-599-413-19	Sequence 19, Appl1	212	29	61.7	515	2	US-10-012-542-219	Sequence 219, App
140	30	63.8	172	2	US-09-599-413-20	Sequence 20, Appl1	213	29	61.7	515	2	US-10-115-123-219	Sequence 219, App
141	30	63.8	172	2	US-10-029-890-2	Sequence 2, Appl1	214	29	61.7	537	2	US-09-388-413B-6	Sequence 6, Appl1
142	30	63.8	195	2	US-09-487-792-11	Sequence 11, Appl1	215	29	61.7	537	2	US-10-233-745-6	Sequence 6, Appl1
143	30	63.8	195	2	US-09-908-594-11	Sequence 11, Appl1	216	29	61.7	666	2	US-09-248-796A-15099	Sequence 15099, A
144	30	63.8	195	2	US-09-487-792-12	Sequence 12, Appl1	217	29	61.7	708	2	US-09-248-796A-15456	Sequence 15456, A
145	30	63.8	196	2	US-09-908-594-12	Sequence 12, Appl1	218	29	61.7	782	2	US-09-585-858-29	Sequence 29, Appl1
146	30	63.8	277	1	US-08-147-784-2	Sequence 2, Appl1	219	29	61.7	834	1	US-10-270-878-29	Sequence 29, Appl1
147	30	63.8	277	2	US-08-195-967-2	Sequence 2, Appl1	220	29	61.7	834	2	US-08-491-357-2	Sequence 2, Appl1
148	30	63.8	277	2	US-09-006-353A-12	Sequence 12, Appl1	221	29	61.7	834	2	US-08-568-633-2	Sequence 2, Appl1
149	30	63.8	277	2	US-08-472-940-2	Sequence 2, Appl1	222	29	61.7	834	2	US-09-196-466-2	Sequence 2, Appl1
150	30	63.8	277	2	US-09-573-986-12	Sequence 12, Appl1	223	29	61.7	834	2	US-09-669-459A-2	Sequence 2, Appl1
151	30	63.8	277	2	US-09-880-939-2	Sequence 2, Appl1	224	29	61.7	834	4	PCT-US96-10823-2	Sequence 2, Appl1
152	30	63.8	277	2	US-09-804-200-2	Sequence 2, Appl1	225	29	61.7	871	4	US-09-303-518B-880	Sequence 880, App
153	30	63.8	277	2	US-08-469-633A-4	Sequence 4, Appl1	226	29	61.7	888	2	US-09-303-518B-878	Sequence 878, App
154	30	63.8	277	2	US-10-046-433-3	Sequence 3, Appl1	227	29	61.7	922	2	US-09-303-518B-882	Sequence 882, App
155	30	63.8	486	2	US-09-734-237B-54	Sequence 54, Appl1	228	29	61.7	922	2	US-09-303-518B-884	Sequence 884, App
156	30	63.8	487	2	US-09-734-237B-56	Sequence 56, Appl1	229	29	61.7	959	2	US-09-314-259-67	Sequence 67, Appl1
157	30	63.8	558	2	US-09-252-991A-21192	Sequence 21192, A	230	29	61.7	961	2	US-09-314-259-66	Sequence 66, Appl1
158	30	63.8	594	2	US-09-949-016-7669	Sequence 7669, Ap	231	29	61.7	1127	2	US-09-150-460B-11	Sequence 11, Appl1
159	30	63.8	604	2	US-09-605-703B-2158	Sequence 2158, Ap	232	29	61.7	1151	2	US-09-328-352-4744	Sequence 4744, Ap
160	30	63.8	681	2	US-09-949-016-5377	Sequence 6377, Ap	233	29	61.7	1650	2	US-09-949-016-11887	Sequence 11887, A
161	30	63.8	705	2	US-09-252-991A-21897	Sequence 21897, A	234	28	59.6	48	2	US-09-902-540-12307	Sequence 12307, A
162	30	63.8	808	1	US-08-629-391A-33	Sequence 33, Appl1	235	28	59.6	88	2	US-09-687-937B-45	Sequence 45, Appl1
163	30	63.8	808	1	US-08-658-335B-33	Sequence 33, Appl1	236	28	59.6	90	2	US-09-352-991A-21264	Sequence 21264, A
164	30	63.8	808	1	US-09-406-640-53	Sequence 33, Appl1	237	28	59.6	115	2	US-09-556-818-1	Sequence 1, Appl1
165	30	63.8	819	1	US-08-464-517-7	Sequence 7, Appl1	238	28	59.6	115	2	US-09-609-027B-6	Sequence 6, Appl1
166	30	63.8	819	1	US-08-464-517-7	Sequence 7, Appl1	239	28	59.6	118	2	US-10-104-047-3683	Sequence 3683, Ap
167	30	63.8	819	4	US-08-463-772-7	Sequence 7, Appl1	240	28	59.6	121	2	US-09-180-864-2	Sequence 2, Appl1
168	30	63.8	819	4	PCT-US93-05000-7	Sequence 7, Appl1	241	28	59.6	124	2	US-09-556-818-4	Sequence 4, Appl1
169	30	63.8	1005	2	US-09-134-000C-6332	Sequence 6322, Ap	242	28	59.6	124	2	US-09-556-818-34	Sequence 34, Appl1
170	30	63.8	1009	2	US-09-762-724-10	Sequence 10, Appl1	243	28	59.6	124	2	US-09-556-818-38	Sequence 38, Appl1
171	30	63.8	1612	4	US-08-545-860D-48	Sequence 48, Appl1	244	28	59.6	126	2	US-09-270-767-37952	Sequence 37952, A
172	30	63.8	1612	4	PCT-US94-04496-48	Sequence 48, Appl1	245	28	59.6	126	2	US-09-270-767-37952	Sequence 37952, A
173	30	63.8	1829	2	US-09-157-420-1	Sequence 1, Appl1	246	28	59.6	126	2	US-09-270-767-53169	Sequence 53169, A

247	28	59.6	126	2	US-09-556-818-2	Sequence 2, Appl1	320	28	59.6	521	2	US-09-270-767-44481	Sequence 44481, A
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249	28	59.6	126	2	US-09-556-818-28	Sequence 28, Appl1	322	28	59.6	627	2	US-09-252-991A-32694	Sequence 32694, A
250	28	59.6	130	2	US-09-556-818-9	Sequence 9, Appl1	323	28	59.6	655	2	US-09-270-767-43477	Sequence 43477, A
251	28	59.6	130	2	US-09-556-818-32	Sequence 32, Appl1	324	28	59.6	671	2	US-09-949-016-9679	Sequence 9679, Ap
252	28	59.6	130	2	US-09-556-818-42	Sequence 42, Appl1	325	28	59.6	765	1	US-08-825-061-19	Sequence 19, Appl1
253	28	59.6	132	2	US-09-556-818-7	Sequence 7, Appl1	326	28	59.6	765	2	US-08-825-886-19	Sequence 19, Appl1
254	28	59.6	132	2	US-09-556-818-10	Sequence 10, Appl1	327	28	59.6	765	2	US-08-989-890-19	Sequence 19, Appl1
255	28	59.6	132	2	US-09-556-818-40	Sequence 40, Appl1	328	28	59.6	878	2	US-09-949-016-10782	Sequence 10782, A
256	28	59.6	134	1	US-08-284-9338-13	Sequence 13, Appl1	329	28	59.6	900	1	US-08-825-061-20	Sequence 20, Appl1
257	28	59.6	134	2	US-08-759-628-9	Sequence 9, Appl1	330	28	59.6	900	1	US-08-825-886-20	Sequence 20, Appl1
258	28	59.6	134	2	US-09-371-615A-7	Sequence 7, Appl1	331	28	59.6	900	2	US-08-989-890-20	Sequence 20, Appl1
259	28	59.6	134	2	US-09-462-941-12	Sequence 12, Appl1	332	28	59.6	914	1	US-08-825-061-21	Sequence 21, Appl1
260	28	59.6	134	4	US-09-462-941-12	Sequence 13, Appl1	333	28	59.6	914	1	US-08-825-886-21	Sequence 21, Appl1
261	28	59.6	137	6	PCT-US95-08950-13	Patent No. 5324640	334	28	59.6	914	2	US-08-989-890-21	Sequence 21, Appl1
262	28	59.6	137	2	US-09-270-767-32781	Sequence 32781, A	335	28	59.6	968	2	US-09-949-016-11229	Sequence 11229, A
263	28	59.6	141	2	US-09-556-818-11	Sequence 11, Appl1	336	28	59.6	1005	2	US-09-252-991A-24655	Sequence 24655, A
264	28	59.6	147	2	US-09-556-818-44	Sequence 44, Appl1	337	28	59.6	1173	2	US-09-543-681A-7965	Sequence 7965, Ap
265	28	59.6	174	2	US-08-706-945D-136	Sequence 136, App	338	28	59.6	1173	2	US-09-248-796A-19313	Sequence 19313, A
266	28	59.6	178	2	US-09-122-443-15	Sequence 15, Appl1	339	28	59.6	1202	1	US-08-825-886-22	Sequence 22, Appl1
267	28	59.6	178	2	US-09-558-089-15	Sequence 15, Appl1	340	28	59.6	1202	1	US-08-825-886-22	Sequence 22, Appl1
268	28	59.6	178	2	US-09-558-087-15	Sequence 15, Appl1	341	28	59.6	1225	2	US-09-949-016-9468	Sequence 9468, Ap
269	28	59.6	178	2	US-09-558-474-15	Sequence 15, Appl1	342	28	59.6	1363	1	US-08-825-886-23	Sequence 23, Appl1
270	28	59.6	190	2	US-09-902-540-15188	Sequence 15188, A	343	28	59.6	1363	1	US-08-825-886-23	Sequence 23, Appl1
271	28	59.6	208	2	US-08-577-788C-50	Sequence 50, Appl1	344	28	59.6	1363	2	US-08-989-890-23	Sequence 23, Appl1
272	28	59.6	228	2	US-09-438-185A-131	Sequence 131, App	345	28	59.6	1436	2	US-09-949-016-6736	Sequence 6736, Ap
273	28	59.6	243	2	US-08-858-207A-338	Sequence 338, App	346	28	59.6	1456	2	US-09-949-016-6795	Sequence 6795, Ap
274	28	59.6	243	2	US-09-198-452A-586	Sequence 586, App	347	28	59.6	1624	2	US-09-949-016-6794	Sequence 6794, Ap
275	28	59.6	247	2	US-09-949-016-8617	Sequence 8617, Ap	348	28	59.6	1676	2	US-09-949-016-7610	Sequence 7610, Ap
276	28	59.6	254	2	US-09-248-796A-2184	Sequence 2184, A	349	28	59.6	1821	2	US-09-949-016-11463	Sequence 11463, A
277	28	59.6	317	2	US-09-270-767-42366	Sequence 42366, A	350	28	59.6	1821	2	US-09-949-016-11465	Sequence 11465, A
278	28	59.6	325	2	US-09-134-000C-6115	Sequence 6115, Ap	351	28	59.6	1821	2	US-09-949-016-11465	Sequence 11465, A
279	28	59.6	330	2	US-09-902-540-10987	Sequence 10987, A	352	28	59.6	1821	2	US-09-949-016-11466	Sequence 11466, A
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281	28	59.6	332	2	US-08-778-394-4	Sequence 4, Appl1	354	28	59.6	1852	1	US-08-825-886-24	Sequence 24, Appl1
282	28	59.6	332	2	US-08-778-423A-2	Sequence 2, Appl1	355	28	59.6	1852	2	US-08-989-890-24	Sequence 24, Appl1
283	28	59.6	344	2	US-09-583-110-3959	Sequence 3959, App	356	28	59.6	1853	1	US-08-425-061-16	Sequence 16, Appl1
284	28	59.6	348	2	US-09-438-185A-550	Sequence 550, App	357	28	59.6	1853	1	US-08-598-591-2	Sequence 2, Appl1
285	28	59.6	356	2	US-09-252-991A-21259	Sequence 21259, A	358	28	59.6	1853	1	US-08-480-784-2	Sequence 2, Appl1
286	28	59.6	359	2	US-09-107-433-6646	Sequence 2646, Ap	359	28	59.6	1853	1	US-08-483-553-2	Sequence 2, Appl1
287	28	59.6	374	2	US-09-949-016-7363	Sequence 20053, Ap	360	28	59.6	1853	1	US-08-487-002-2	Sequence 2, Appl1
288	28	59.6	392	2	US-09-252-991A-20053	Sequence 12088, A	361	28	59.6	1853	1	US-08-483-554B-2	Sequence 2, Appl1
289	28	59.6	394	2	US-09-489-039A-12088	Sequence 147, App	362	28	59.6	1853	1	US-08-798-691-2	Sequence 4, Appl1
290	28	59.6	396	2	US-09-198-452A-147	Sequence 228, App	363	28	59.6	1853	1	US-08-798-691-4	Sequence 6, Appl1
291	28	59.6	398	2	US-09-198-452A-228	Sequence 214, App	364	28	59.6	1853	1	US-08-798-691-6	Sequence 2, Appl1
292	28	59.6	398	2	US-09-438-185A-214	Sequence 2, Appl1	365	28	59.6	1853	1	US-08-825-886-16	Sequence 16, Appl1
293	28	59.6	401	2	US-08-974-022-2	Sequence 3, Appl1	366	28	59.6	1853	1	US-08-825-886-16	Sequence 2, Appl1
294	28	59.6	401	2	US-08-289-222E-3	Sequence 3, Appl1	367	28	59.6	1853	1	US-08-825-886-16	Sequence 2, Appl1
295	28	59.6	401	2	US-09-054-526B-3	Sequence 2, Appl1	368	28	59.6	1853	1	US-08-825-886-16	Sequence 2, Appl1
296	28	59.6	401	2	US-08-795-445A-2	Sequence 2, Appl1	369	28	59.6	1853	1	US-08-825-886-16	Sequence 2, Appl1
297	28	59.6	401	2	US-08-795-447A-2	Sequence 2, Appl1	370	28	59.6	1853	1	US-08-825-886-16	Sequence 2, Appl1
298	28	59.6	401	2	US-08-974-186-2	Sequence 2, Appl1	371	28	59.6	1853	1	US-09-074-476-2	Sequence 2, Appl1
299	28	59.6	401	2	US-08-795-446B-2	Sequence 2, Appl1	372	28	59.6	1853	1	US-09-074-476-4	Sequence 4, Appl1
300	28	59.6	401	2	US-08-706-945D-124	Sequence 124, App	373	28	59.6	1853	1	US-09-074-476-6	Sequence 6, Appl1
301	28	59.6	401	2	US-08-577-788C-2	Sequence 2, Appl1	374	28	59.6	1853	1	US-09-099-753-2	Sequence 2, Appl1
302	28	59.6	401	2	US-08-577-788C-55	Sequence 55, Appl1	375	28	59.6	1853	1	US-08-850-727-2	Sequence 2, Appl1
303	28	59.6	411	1	US-08-310-693-2	Sequence 2, Appl1	376	28	59.6	1853	1	US-08-850-727-2	Sequence 2, Appl1
304	28	59.6	411	4	PCT-US95-11280-2	Sequence 2, Appl1	377	28	59.6	1853	1	US-08-907-678B-49	Sequence 49, Appl1
305	28	59.6	423	2	US-09-674-330A-4	Sequence 4, Appl1	378	28	59.6	1853	1	US-08-989-890-16	Sequence 16, Appl1
306	28	59.6	423	2	US-09-674-330A-9	Sequence 9, Appl1	379	28	59.6	1853	1	US-10-022-819-2	Sequence 2, Appl1
307	28	59.6	442	2	US-09-540-236-2820	Sequence 2820, Ap	380	28	59.6	1853	1	US-09-462-01A-2	Sequence 1098, Ap
308	28	59.6	442	2	US-09-674-330A-3	Sequence 3, Appl1	381	28	59.6	1853	1	US-09-734-672A-2	Sequence 2, Appl1
309	28	59.6	454	2	US-09-252-991A-28000	Sequence 28000, A	382	28	59.6	1853	1	US-09-734-672A-4	Sequence 4, Appl1
310	28	59.6	461	2	US-09-674-330A-8	Sequence 8, Appl1	383	28	59.6	1853	1	US-09-734-672A-6	Sequence 6, Appl1
311	28	59.6	484	2	US-09-902-540-13700	Sequence 13700, A	384	28	59.6	1853	1	US-09-923-327A-264	Sequence 264, App
312	28	59.6	495	1	US-08-455-559-10	Sequence 10, Appl1	385	28	59.6	1853	1	US-09-923-327A-266	Sequence 266, App
313	28	59.6	495	1	US-09-145-060-10	Sequence 10, Appl1	386	28	59.6	1853	1	US-09-923-327A-268	Sequence 268, App
314	28	59.6	495	4	PCT-US94-00657-10	Sequence 10, Appl1	387	28	59.6	1853	1	PCT-US95-10202-2	Sequence 2, Appl1
315	28	59.6	498	2	US-09-328-352-7983	Sequence 7983, Ap	388	28	59.6	1853	1	PCT-US95-10203-2	Sequence 2, Appl1
316	28	59.6	501	1	US-08-288-508C-2	Sequence 2, Appl1	389	28	59.6	1853	1	PCT-US95-10220-2	Sequence 2, Appl1
317	28	59.6	501	2	US-08-981-490B-1	Sequence 1, Appl1	390	28	59.6	1853	1	PCT-US95-10220-2	Sequence 2, Appl1
318	28	59.6	501	2	US-09-386-450D-2	Sequence 2, Appl1	391	28	59.6	1853	4	PCT-US95-10220-2	Sequence 2, Appl1
319	28	59.6	501	2	US-09-949-016-5956	Sequence 5956, Ap	392	28	59.6	2210	2	US-09-309-572-7	Sequence 7, Appl1

393	28	59.6	2210	2	US-09-718-096-7	Sequence 7, Appl1	466	27	57.4	167	2	US-09-489-039A-12336	Sequence 12336, A
394	28	59.6	3729	1	US-08-804-227C-4	Sequence 4, Appl1	467	27	57.4	168	2	US-09-602-777A-196	Sequence 196, Appl
395	28	59.6	4861	2	US-09-919-497-70	Sequence 70, Appl	468	27	57.4	170	2	US-09-270-767-53064	Sequence 53064, A
396	27	57.4	10	2	US-08-159-339A-559	Sequence 559, Appl	469	27	57.4	184	2	US-09-570-856B-25	Sequence 25, Appl1
397	27	57.4	11	2	US-09-601-729-176	Sequence 176, App	470	27	57.4	185	2	US-09-230-590-2	Sequence 2, Appl1
398	27	57.4	17	6	5336758-1	Patent No. 5336758	471	27	57.4	185	2	US-09-970-616-2	Sequence 2, Appl1
399	27	57.4	21	2	US-09-142-569-17	Sequence 17, Appl1	472	27	57.4	201	2	US-09-248-796A-77602	Sequence 27602, A
400	27	57.4	21	2	US-09-495-448A-17	Sequence 17, Appl1	473	27	57.4	210	2	US-09-134-000C-6797	Sequence 6797, Ap
401	27	57.4	22	2	US-08-851-843A-160	Sequence 160, App	474	27	57.4	211	1	US-07-842-349-2	Sequence 2, Appl1
402	27	57.4	22	2	US-08-974-549A-280	Sequence 280, App	475	27	57.4	211	1	US-08-244-686-2	Sequence 2, Appl1
403	27	57.4	22	2	US-08-854-050-160	Sequence 160, App	476	27	57.4	211	2	US-07-955-726A-8	Sequence 8, Appl1
404	27	57.4	22	2	US-09-430-323-160	Sequence 160, App	477	27	57.4	211	2	US-09-570-856B-27	Sequence 27, Appl1
405	27	57.4	22	2	US-09-402-181B-280	Sequence 280, App	478	27	57.4	211	2	US-09-570-856B-33	Sequence 33, Appl1
406	27	57.4	22	2	US-09-721-456-280	Sequence 280, App	479	27	57.4	213	2	US-09-949-016-9284	Sequence 9284, Ap
407	27	57.4	22	2	US-09-766-253-160	Sequence 160, App	480	27	57.4	221	2	US-09-270-767-47505	Sequence 47505, A
408	27	57.4	22	2	US-10-054-295-160	Sequence 160, App	481	27	57.4	224	2	US-09-270-767-43678	Sequence 43678, A
409	27	57.4	22	2	US-09-438-486A-160	Sequence 160, App	482	27	57.4	226	2	US-09-583-110-3394	Sequence 3394, Ap
410	27	57.4	34	2	US-08-851-843A-138	Sequence 138, App	483	27	57.4	229	2	US-09-247-155-97	Sequence 97, Appl
411	27	57.4	34	2	US-08-974-549A-256	Sequence 256, App	484	27	57.4	229	2	US-09-991-181-474	Sequence 424, App
412	27	57.4	34	2	US-08-854-050-138	Sequence 138, App	485	27	57.4	229	2	US-09-990-444-424	Sequence 424, App
413	27	57.4	34	2	US-09-430-323-138	Sequence 138, App	486	27	57.4	229	2	US-09-903-190-97	Sequence 97, Appl
414	27	57.4	34	2	US-09-402-181B-256	Sequence 256, App	487	27	57.4	229	2	US-09-997-333-424	Sequence 424, App
415	27	57.4	34	2	US-09-721-456-256	Sequence 256, App	488	27	57.4	229	2	US-09-992-598-424	Sequence 424, App
416	27	57.4	34	2	US-09-766-253-138	Sequence 138, App	489	27	57.4	233	2	US-08-851-843A-64	Sequence 64, Appl
417	27	57.4	34	2	US-10-054-295-138	Sequence 138, App	490	27	57.4	233	2	US-08-974-549A-15	Sequence 15, Appl
418	27	57.4	34	2	US-09-438-486A-138	Sequence 138, App	491	27	57.4	233	2	US-08-854-050-64	Sequence 64, Appl
419	27	57.4	42	2	US-08-974-549A-96	Sequence 96, Appl1	492	27	57.4	233	2	US-09-430-323-64	Sequence 64, Appl
420	27	57.4	42	2	US-08-912-951-96	Sequence 96, Appl1	493	27	57.4	233	2	US-09-912-951-15	Sequence 15, Appl
421	27	57.4	42	2	US-09-402-181B-96	Sequence 96, Appl1	494	27	57.4	233	2	US-09-328-352-6789	Sequence 6789, Ap
422	27	57.4	42	2	US-09-721-456-96	Sequence 96, Appl1	495	27	57.4	233	2	US-09-402-181B-15	Sequence 15, Appl1
423	27	57.4	47	2	US-08-974-549A-42	Sequence 42, Appl	496	27	57.4	233	2	US-09-721-456-15	Sequence 15, Appl
424	27	57.4	47	2	US-08-912-951-42	Sequence 42, Appl	497	27	57.4	233	2	US-09-766-253-64	Sequence 64, Appl
425	27	57.4	47	2	US-09-402-181B-42	Sequence 42, Appl	498	27	57.4	233	2	US-10-054-295-64	Sequence 64, Appl
426	27	57.4	47	2	US-09-721-456-42	Sequence 42, Appl	499	27	57.4	233	2	US-09-438-486A-73	Sequence 73, Appl
427	27	57.4	47	2	US-09-442-460-43	Sequence 43, Appl1	500	27	57.4	237	2	US-08-679-493A-73	Sequence 73, Appl
428	27	57.4	49	1	US-08-417-210A-114	Sequence 114, App	501	27	57.4	251	2	US-09-489-039A-10609	Sequence 10609, A
429	27	57.4	49	2	US-09-136-159A-114	Sequence 114, App	502	27	57.4	251	2	US-09-107-532A-4381	Sequence 4381, Ap
430	27	57.4	60	2	US-10-179-784-12	Sequence 12, Appl	503	27	57.4	253	2	US-09-270-767-33689	Sequence 33689, A
431	27	57.4	61	2	US-09-270-767-40422	Sequence 40422, A	504	27	57.4	258	2	US-09-540-236-2125	Sequence 2125, Ap
432	27	57.4	61	2	US-09-270-767-55638	Sequence 55638, A	505	27	57.4	261	6	5320958-3	Patent No. 5320958
433	27	57.4	67	2	US-09-248-796A-24808	Sequence 24808, A	506	27	57.4	263	1	US-08-362-670B-32	Sequence 32, Appl
434	27	57.4	70	2	US-09-448-796A-28035	Sequence 28035, A	507	27	57.4	263	2	US-08-333-576C-32	Sequence 32, Appl
435	27	57.4	76	2	US-09-270-767-41257	Sequence 41257, A	508	27	57.4	263	2	US-08-808-324-32	Sequence 32, Appl
436	27	57.4	76	2	US-09-270-767-56473	Sequence 56473, A	509	27	57.4	263	2	US-09-945-182-32	Sequence 32, Appl
437	27	57.4	83	2	US-09-762-724-15	Sequence 16, Appl	510	27	57.4	263	4	PCT-US94-14030A-32	Sequence 32, Appl
438	27	57.4	84	2	US-09-270-767-32880	Sequence 32880, A	511	27	57.4	266	2	PCT-US94-14030A-32	Sequence 32, Appl
439	27	57.4	84	2	US-09-270-767-48097	Sequence 48097, A	512	27	57.4	271	1	US-08-956-809-5	Sequence 5, Appl1
440	27	57.4	86	2	US-09-252-991A-16624	Sequence 16624, A	513	27	57.4	273	2	US-09-248-796A-16972	Sequence 16972, A
441	27	57.4	91	2	US-09-270-767-34473	Sequence 34473, A	514	27	57.4	274	2	US-09-252-991A-23302	Sequence 23302, A
442	27	57.4	91	2	US-09-270-767-49690	Sequence 49690, A	515	27	57.4	275	6	5252477-3	Patent No. 5252477
443	27	57.4	107	6	5514582-17	Patent No. 5514582	516	27	57.4	280	1	US-08-816-155B-43	Sequence 43, Appl
444	27	57.4	108	6	5514582-16	Patent No. 5514582	517	27	57.4	280	2	US-09-079-587-43	Sequence 43, Appl
445	27	57.4	123	2	US-09-328-352-6963	Sequence 6963, Ap	518	27	57.4	281	2	US-09-252-991A-25471	Sequence 25471, A
446	27	57.4	137	2	US-09-270-767-32748	Sequence 32748, A	519	27	57.4	284	2	US-09-302-626B-58	Sequence 58, Appl
447	27	57.4	137	2	US-09-270-767-47965	Sequence 47965, A	520	27	57.4	292	2	US-09-134-001C-3515	Sequence 3515, Ap
448	27	57.4	138	2	US-09-489-039A-7633	Sequence 7633, Ap	521	27	57.4	293	2	US-09-489-039A-11514	Sequence 11514, A
449	27	57.4	141	2	US-09-270-767-33783	Sequence 33783, A	522	27	57.4	300	2	US-09-270-767-34413	Sequence 34413, A
450	27	57.4	141	2	US-09-270-767-49000	Sequence 49000, A	523	27	57.4	300	2	US-09-270-767-49630	Sequence 49630, A
451	27	57.4	144	2	US-09-270-767-41052	Sequence 41052, A	524	27	57.4	308	2	US-09-369-247-60	Sequence 60, Appl
452	27	57.4	144	2	US-09-270-767-56268	Sequence 56268, A	525	27	57.4	308	2	US-10-062-548-60	Sequence 60, Appl
453	27	57.4	149	2	US-09-270-767-33323	Sequence 33323, A	526	27	57.4	311	2	US-09-543-681A-5602	Sequence 5602, Ap
454	27	57.4	149	2	US-09-270-767-48540	Sequence 48540, A	527	27	57.4	314	1	US-08-589-446-6	Sequence 6, Appl1
455	27	57.4	157	2	US-08-851-843A-83	Sequence 83, Appl	528	27	57.4	314	1	US-08-444-882-6	Sequence 6, Appl1
456	27	57.4	157	2	US-08-974-549A-266	Sequence 226, App	529	27	57.4	314	1	US-08-389-459A-6	Sequence 6, Appl1
457	27	57.4	157	2	US-08-854-050-83	Sequence 83, Appl	530	27	57.4	314	2	US-08-987-8678A-26	Sequence 26, Appl1
458	27	57.4	157	2	US-09-430-323-83	Sequence 83, Appl	531	27	57.4	321	1	US-08-362-670B-26	Sequence 26, Appl1
459	27	57.4	157	2	US-09-402-181B-226	Sequence 226, App	532	27	57.4	321	2	US-08-333-576C-26	Sequence 26, Appl
460	27	57.4	157	2	US-09-721-456-226	Sequence 226, App	533	27	57.4	321	2	US-08-808-324-26	Sequence 26, Appl
461	27	57.4	157	2	US-09-766-253-83	Sequence 83, Appl	534	27	57.4	321	2	US-09-945-182-26	Sequence 26, Appl
462	27	57.4	157	2	US-10-054-295-83	Sequence 83, Appl	535	27	57.4	321	2	PCT-US94-14030A-26	Sequence 26, Appl
463	27	57.4	157	2	US-09-438-486A-83	Sequence 83, Appl	536	27	57.4	327	2	US-08-679-493A-71	Sequence 71, Appl
464	27	57.4	162	2	US-09-270-767-40554	Sequence 40554, A	537	27	57.4	327	2	US-08-679-493A-72	Sequence 72, Appl
465	27	57.4	162	2	US-09-270-767-55770	Sequence 55770, A	538	27	57.4	327	2	US-10-104-047-2508	Sequence 2508, Ap

539	27	57.4	335	2	US-09-312-283C-186	Sequence 186, App	612	27	57.4	892	2	US-10-226-629A-15	Sequence 15, Appl
540	27	57.4	337	2	US-09-188-930-186	Sequence 186, App	613	27	57.4	913	1	US-07-743-357-6	Sequence 6, Appl
541	27	57.4	342	2	US-09-136-073-2	Sequence 2, Appl	614	27	57.4	913	1	US-07-743-357-22	Sequence 22, Appl
542	27	57.4	342	2	US-09-457-024A-2	Sequence 2, Appl	615	27	57.4	936	2	US-10-104-047-2621	Sequence 2621, Ap
543	27	57.4	360	2	US-09-302-626B-178	Sequence 178, App	616	27	57.4	1000	2	US-10-290-579A-186	Sequence 186, App
544	27	57.4	364	2	US-09-417-485D-46	Sequence 46, Appl	617	27	57.4	1002	2	US-10-290-579A-185	Sequence 185, App
545	27	57.4	379	1	US-08-468-847B-11	Sequence 11, Appl	618	27	57.4	1003	1	US-07-743-357-8	Sequence 8, Appl
546	27	57.4	379	2	US-09-142-569-2	Sequence 2, Appl	619	27	57.4	1003	2	US-09-309-572-17	Sequence 17, Appl
547	27	57.4	379	2	US-09-495-448A-2	Sequence 2, Appl	620	27	57.4	1003	2	US-09-718-096-17	Sequence 17, Appl
548	27	57.4	397	2	US-09-270-767-41811	Sequence 41811, A	621	27	57.4	1003	2	US-10-290-579A-188	Sequence 188, App
549	27	57.4	399	1	US-09-414-926A-5	Sequence 5, Appl	622	27	57.4	1003	2	US-09-333-795-7	Sequence 7, Appl
550	27	57.4	399	1	US-08-926-922-5	Sequence 5, Appl	623	27	57.4	1003	2	US-08-463-795-9	Sequence 9, Appl
551	27	57.4	399	2	US-09-253-682-5	Sequence 5, Appl	624	27	57.4	1003	2	US-09-393-795-9	Sequence 11, Appl
552	27	57.4	399	2	US-09-527-657-5	Sequence 5, Appl	625	27	57.4	1004	2	US-09-393-795-11	Sequence 7, Appl
553	27	57.4	405	2	US-09-892-100-5	Sequence 5, Appl	626	27	57.4	1005	1	US-07-743-357-7	Sequence 7, Appl
554	27	57.4	405	2	US-09-489-039A-9126	Sequence 9126, Ap	627	27	57.4	1014	2	US-09-319-588C-6	Sequence 6, Appl
555	27	57.4	408	2	US-09-315-793-32	Sequence 32, Appl	628	27	57.4	1015	2	US-08-463-795-9	Sequence 9, Appl
556	27	57.4	427	2	US-09-690-265-1	Sequence 1, Appl	629	27	57.4	1015	2	US-09-124-900-3	Sequence 3, Appl
557	27	57.4	427	2	US-09-205-785-1	Sequence 1, Appl	630	27	57.4	1015	2	US-08-463-028-9	Sequence 9, Appl
558	27	57.4	435	2	US-10-179-784-2	Sequence 2, Appl	631	27	57.4	1015	2	US-08-463-209-9	Sequence 9, Appl
559	27	57.4	438	2	US-09-252-991A-19934	Sequence 19934, A	632	27	57.4	1016	1	US-07-743-357-2	Sequence 2, Appl
560	27	57.4	446	2	US-09-270-767-33198	Sequence 33198, A	633	27	57.4	1016	1	US-07-743-357-3	Sequence 3, Appl
561	27	57.4	446	2	US-09-270-767-48415	Sequence 48415, A	634	27	57.4	1016	1	US-07-743-357-4	Sequence 4, Appl
562	27	57.4	446	2	US-09-902-540-11040	Sequence 11040, A	635	27	57.4	1016	1	US-07-743-357-5	Sequence 5, Appl
563	27	57.4	469	2	US-09-248-796A-17534	Sequence 17534, A	636	27	57.4	1017	2	US-09-762-724-12	Sequence 12, Appl
564	27	57.4	481	2	US-09-673-395A-277	Sequence 277, App	637	27	57.4	1023	2	US-09-762-724-14	Sequence 14, Appl
565	27	57.4	484	2	US-09-270-767-44606	Sequence 44606, A	638	27	57.4	1027	2	US-09-762-724-6	Sequence 6, Appl
566	27	57.4	489	2	US-10-104-047-2922	Sequence 2922, Ap	639	27	57.4	1029	2	US-09-762-724-8	Sequence 8, Appl
567	27	57.4	496	2	US-09-171-965-2	Sequence 2, Appl	640	27	57.4	1151	2	US-08-836-134-23	Sequence 23, Appl
568	27	57.4	496	2	US-08-945-289-6	Sequence 6, Appl	641	27	57.4	1151	2	US-09-493-784-23	Sequence 23, Appl
569	27	57.4	496	2	US-09-845-511-2	Sequence 2, Appl	642	27	57.4	1232	2	US-08-836-134-2	Sequence 2, Appl
570	27	57.4	498	2	US-09-232-468A-24	Sequence 24, Appl	643	27	57.4	1232	2	US-09-493-784-2	Sequence 2, Appl
571	27	57.4	498	2	US-09-784-984B-54	Sequence 54, Appl	644	27	57.4	1276	2	US-09-949-016-8886	Sequence 8886, Ap
572	27	57.4	513	2	US-09-949-016-5900	Sequence 5900, Ap	645	27	57.4	1295	2	US-09-705-872-3	Sequence 3, Appl
573	27	57.4	515	1	US-08-063-552-4	Sequence 4, Appl	646	27	57.4	1350	2	US-09-952-060-35	Sequence 35, Appl
574	27	57.4	515	4	PCT-US93-05704-4	Sequence 4, Appl	647	27	57.4	1388	1	US-08-685-576-1	Sequence 1, Appl
575	27	57.4	519	2	US-09-248-796A-19263	Sequence 19263, A	648	27	57.4	1403	2	US-09-705-872-1	Sequence 1, Appl
576	27	57.4	542	2	US-09-248-796A-15128	Sequence 15128, A	649	27	57.4	1403	2	US-09-949-002-370	Sequence 370, App
577	27	57.4	562	2	US-09-117-217-14	Sequence 14, Appl	650	27	57.4	1412	2	US-09-949-002-411	Sequence 411, App
578	27	57.4	562	2	US-09-335-487-14	Sequence 14, Appl	651	27	57.4	1564	2	US-09-302-626B-62	Sequence 62, App
579	27	57.4	579	2	US-09-198-452A-918	Sequence 918, App	652	27	57.4	1574	2	US-09-487-558B-244	Sequence 244, App
580	27	57.4	602	2	US-09-902-540-16677	Sequence 16677, A	653	27	57.4	1648	2	US-09-302-626B-179	Sequence 179, App
581	27	57.4	606	2	US-09-789-272-2	Sequence 2, Appl	654	27	57.4	1711	2	US-09-515-806-4	Sequence 4, Appl
582	27	57.4	636	2	US-09-438-185A-853	Sequence 853, App	655	27	57.4	1711	2	US-09-771-161A-219	Sequence 219, App
583	27	57.4	633	2	US-09-328-352-6519	Sequence 6519, App	656	27	57.4	1978	2	US-09-771-161A-220	Sequence 220, App
584	27	57.4	645	2	US-09-902-540-16286	Sequence 16286, A	657	27	57.4	2047	2	US-09-302-626B-60	Sequence 60, App
585	27	57.4	655	2	US-10-104-047-2502	Sequence 2502, Ap	658	27	57.4	3542	2	US-09-949-016-6468	Sequence 6468, Ap
586	27	57.4	660	2	US-09-270-767-46777	Sequence 46777, A	659	27	57.4	3542	2	US-09-949-016-7404	Sequence 7404, Ap
587	27	57.4	671	2	US-09-708-426-11	Sequence 11, Appl	660	27	57.4	8	2	US-10-087-013-2	Sequence 2, Appl
588	27	57.4	674	2	US-09-489-039A-9158	Sequence 9158, Ap	661	27	57.4	8	2	US-09-314-268-65	Sequence 65, Appl
589	27	57.4	712	2	US-08-872-855-9	Sequence 9, Appl	662	27	57.4	14	2	US-10-394-980-55	Sequence 55, Appl
590	27	57.4	734	2	US-09-949-016-6415	Sequence 6415, Ap	663	27	57.4	14	2	US-09-314-268-66	Sequence 66, Appl
591	27	57.4	766	1	US-08-846-762-5	Sequence 5, Appl	664	27	57.4	20	1	US-09-949-002-411	Sequence 411, App
592	27	57.4	800	2	US-09-489-039A-7191	Sequence 7191, Ap	665	27	57.4	20	1	US-08-218-025A-10	Sequence 10, Appl
593	27	57.4	818	2	US-09-248-796A-20792	Sequence 20792, A	666	27	57.4	24	2	US-08-934-915-172	Sequence 172, App
594	27	57.4	850	2	US-09-904-389-2	Sequence 2, Appl	667	27	57.4	26	2	US-09-461-697-177	Sequence 177, App
595	27	57.4	850	2	US-09-952-060-2	Sequence 2, Appl	668	27	57.4	30	1	US-09-842-164A-38	Sequence 38, Appl
596	27	57.4	850	2	US-09-952-060-4	Sequence 4, Appl	669	27	57.4	30	1	US-08-934-915-151	Sequence 151, Appl
597	27	57.4	869	2	US-09-540-236-2799	Sequence 2799, Ap	670	27	57.4	62	2	US-09-248-796A-6881	Sequence 6881, A
598	27	57.4	875	2	US-09-952-060-6	Sequence 6, Appl	671	27	57.4	66	2	US-09-270-767-61766	Sequence 61766, A
599	27	57.4	884	2	US-08-851-843A-55	Sequence 55, Appl	672	27	57.4	75	2	US-09-248-796A-22537	Sequence 22537, A
600	27	57.4	884	2	US-08-974-549A-222	Sequence 222, Appl	673	27	57.4	76	2	US-09-540-236-2179	Sequence 2179, Ap
601	27	57.4	884	2	US-08-854-050-55	Sequence 55, Appl	674	27	57.4	76	2	US-09-270-767-40176	Sequence 40176, A
602	27	57.4	884	2	US-09-430-323-55	Sequence 55, Appl	675	27	57.4	79	2	US-09-543-681A-7226	Sequence 7226, Ap
603	27	57.4	884	2	US-09-402-181B-222	Sequence 222, App	676	27	57.4	92	2	US-09-314-268-90	Sequence 90, Appl
604	27	57.4	884	2	US-09-721-456-222	Sequence 222, App	677	27	57.4	100	2	US-09-034-916-13	Sequence 13, Appl
605	27	57.4	884	2	US-09-766-253-55	Sequence 55, Appl	678	27	57.4	103	2	US-09-191-885-21	Sequence 21, Appl
606	27	57.4	884	2	US-09-502-498C-5	Sequence 5, Appl	679	27	57.4	103	2	US-08-817-906-21	Sequence 21, Appl
607	27	57.4	884	2	US-09-502-498C-5	Sequence 5, Appl	680	27	57.4	103	4	PCT-US95-13376-21	Sequence 21, Appl
608	27	57.4	884	2	US-09-502-424C-5	Sequence 5, Appl	681	27	57.4	124	2	US-09-902-540-12036	Sequence 12036, A
609	27	57.4	884	2	US-10-054-295-55	Sequence 55, Appl	682	27	57.4	125	2	US-09-328-352-7243	Sequence 7243, Ap
610	27	57.4	884	2	US-09-438-486A-55	Sequence 55, Appl	683	27	57.4	127	1	US-08-162-146-3	Sequence 3, Appl
611	27	57.4	891	2	US-10-226-629A-16	Sequence 16, Appl	684	27	57.4	127	2	US-09-314-127-3	Sequence 3, Appl

685	26	55.3	129	2	US-09-270-767-40305	Sequence 40305, A	758	26	55.3	307	2	US-10-009-962-7	Sequence 7, Appl1
686	26	55.3	129	2	US-09-270-767-55521	Sequence 55521, A	759	26	55.3	307	2	US-10-037-417-87	Sequence 87, Appl1
687	26	55.3	131	1	US-08-721-488-8	Sequence 8, Appl1	760	26	55.3	307	2	US-10-037-417-88	Sequence 88, Appl1
688	26	55.3	131	2	US-09-270-767-40433	Sequence 40433, A	761	26	55.3	307	2	US-10-671-628-7	Sequence 7, Appl1
689	26	55.3	131	2	US-09-270-767-55649	Sequence 55649, A	762	26	55.3	309	2	US-09-949-016-10401	Sequence 10401, A
690	26	55.3	131	2	US-09-538-092-620	Sequence 620, App	763	26	55.3	314	2	US-09-328-352-4934	Sequence 4934, Ap
691	26	55.3	138	2	US-08-914-479A-2	Sequence 2, Appl1	764	26	55.3	320	2	US-09-248-766A-19339	Sequence 19339, A
692	26	55.3	139	2	US-10-104-047-2691	Sequence 2691, Ap	765	26	55.3	320	2	US-09-328-352-7120	Sequence 7120, Ap
693	26	55.3	140	2	US-10-104-047-2719	Sequence 2719, Ap	766	26	55.3	323	2	US-09-949-016-7924	Sequence 7924, Ap
694	26	55.3	140	2	US-09-252-991A-2640	Sequence 26480, A	767	26	55.3	323	2	US-10-104-047-2839	Sequence 2839, Ap
695	26	55.3	149	2	US-09-270-767-41757	Sequence 41757, A	768	26	55.3	338	2	US-09-739-455-9	Sequence 9, Appl1
696	26	55.3	155	2	US-09-270-767-32193	Sequence 32193, A	769	26	55.3	338	2	US-09-739-455-11	Sequence 10, Appl1
697	26	55.3	155	2	US-09-248-796A-20912	Sequence 20912, A	770	26	55.3	338	2	US-09-739-455-11	Sequence 11, Appl1
698	26	55.3	159	2	US-09-134-001C-3314	Sequence 3314, Ap	771	26	55.3	338	2	US-09-739-455-19	Sequence 19, Appl1
699	26	55.3	163	2	US-09-270-767-40798	Sequence 40798, A	772	26	55.3	338	2	US-09-739-455-20	Sequence 20, Appl1
700	26	55.3	163	2	US-09-270-767-56014	Sequence 56014, A	773	26	55.3	338	2	US-09-739-455-21	Sequence 21, Appl1
701	26	55.3	169	2	US-09-270-767-60047	Sequence 60047, A	774	26	55.3	338	2	US-10-153-919-9	Sequence 9, Appl1
702	26	55.3	171	2	US-09-248-796A-21440	Sequence 21420, A	775	26	55.3	338	2	US-10-153-919-10	Sequence 10, Appl1
703	26	55.3	175	2	US-10-101-464A-583	Sequence 583, App	776	26	55.3	338	2	US-10-153-919-11	Sequence 11, Appl1
704	26	55.3	177	2	US-10-104-047-3347	Sequence 3347, Ap	777	26	55.3	338	2	US-10-153-919-19	Sequence 19, Appl1
705	26	55.3	178	2	US-09-495-066-2	Sequence 2, Appl1	778	26	55.3	338	2	US-10-153-919-20	Sequence 20, Appl1
706	26	55.3	178	2	US-09-107-433-4398	Sequence 4398, Ap	779	26	55.3	338	2	US-10-153-919-21	Sequence 21, Appl1
707	26	55.3	180	2	US-09-328-352-8083	Sequence 8083, Ap	780	26	55.3	345	2	US-09-543-681A-4540	Sequence 4540, Ap
708	26	55.3	181	2	US-09-270-767-1838	Sequence 31838, A	781	26	55.3	335	2	US-09-270-767-46203	Sequence 46203, A
709	26	55.3	181	2	US-09-270-767-47055	Sequence 47055, A	782	26	55.3	333	2	US-09-489-039A-13640	Sequence 13640, A
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979 25 53.2 40 2 US-09-673-395A-466
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ALIGNMENTS

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Sequence 12332, A
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Sequence 12, Appl
Sequence 101, App
Sequence 3, Appl1
Sequence 26405, A
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Sequence 26588, A

RESULT 1
; Sequence 159, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
; TITLE OF INVENTION: DIAGNOSTIC PURPOSES
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; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. Foutch
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946,6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:
; INFORMATION FOR SEQ ID NO: 159:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid

TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-934-915-159

Query Match 100.0%; Score 47; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.037;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLTCTELQTT 9
Db 5 QLTCTELQTT 13

RESULT 2
US-08-363-586-4
; Sequence 4, Application US/08363586
; Patent No. 5629161
; GENERAL INFORMATION:
; APPLICANT: Mueller, Martin
; APPLICANT: Giesmann, Lutz
; TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
; TITLE OF INVENTION: Peptides for the Diagnostic Purpose
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; ADDRESSEE: Dunner
; STREET: 1300 I Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/363,586
; FILING DATE: 23-DEC-1994
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/909,296
; FILING DATE: 09-JUL-1992
; APPLICATION NUMBER: EP 91111720.8
; FILING DATE: 13-JUL-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Madler, Linda A.
; REGISTRATION NUMBER: 33,218
; REFERENCE/DOCKET NUMBER: 02481-1195-00000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-408-4000
; TELEFAX: 202-408-4400
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 30 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-363-586-4

Query Match 100.0%; Score 47; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.055;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLTCTELQTT 9
Db 14 QLTCTELQTT 22

RESULT 3
US-09-980-523A-4
; Sequence 4, Application US/09980523A
; Patent No. 6783763

```

; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
; US-09-980-523A-4

Query Match          100.0%; Score 47; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.055;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
DB 7 QLCTELQTT 15

RESULT 4
US-09-701-080C-18
; Sequence 18, Application US/09701080C
; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; PRIOR FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
; US-09-701-080C-18

Query Match          100.0%; Score 47; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.27;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
DB 14 QLCTELQTT 22

RESULT 5
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD

```

```

; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
; US-09-980-523A-2

Query Match          100.0%; Score 47; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.28;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
DB 21 QLCTELQTT 29

RESULT 6
US-08-316-239B-3
; Sequence 3, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
; US-08-316-239B-3

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Query Match 100.0%; Score 47; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.29; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
|||
Db 21 QLCLEQTT 29

RESULT 7

US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESS: Jagtland & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentln Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtland, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
; US-08-316-239B-4

Query Match 100.0%; Score 47; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.29; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
|||
Db 21 QLCLEQTT 29

RESULT 8

US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian

; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentln Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
; US-08-860-165-14

Query Match 100.0%; Score 47; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.31; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
|||
Db 90 QLCLEQTT 98

RESULT 9

US-09-359-382-14
; Sequence 14, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentln Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
; US-09-359-382-14

Query Match 100.0%; Score 47; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.31; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
|||
Db 90 QLCLEQTT 98

RESULT 10

US-08-117-083-10
; Sequence 10, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Bouras, Michael E.
; APPLICANT: Inglis, Stephen C.
; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; TITLE OF INVENTION: Papilloma Virus Proteins

```

;
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Walter H. Dreger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
;
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
;
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
;
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dreger, Walter H.
; REGISTRATION NUMBER: 24,190
; REFERENCE/DOCKET NUMBER: A-58763
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
;
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 182 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
;
; NAME/KEY: Protein
; LOCATION: 1..182
; OTHER INFORMATION: /note="Xaa refers to stop codon in
;
; US-08-117-083-10
;
Query Match 100.0%; Score 47; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
Db 22 QLCTELQTT 30

RESULT 11
US-09-462-993-1
; Sequence 1, Application US/09462993
; Patent No. 6884786
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOU, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Derived from
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```

;
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
;
; OTHER INFORMATION: fused F protein signals, clone E6*TMF.
;
; US-09-462-993-1
;
Query Match 100.0%; Score 47; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.43;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
Db 49 QLCTELQTT 57

RESULT 12
US-08-860-165-10
; Sequence 10, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
;
; US-08-860-165-10
;
Query Match 100.0%; Score 47; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
Db 21 QLCTELQTT 29

RESULT 13
US-09-359-382-10
; Sequence 10, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
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ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 47; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
| | | | | | | | | |
DB 21 QLCTELQTT 29

RESULT 14

US-09-367-309A-1
Sequence 1, Application US/09367309A
Patent No. 6428607
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 47; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
| | | | | | | | | |
DB 21 QLCTELQTT 29

RESULT 15

US-09-485-885-4
Sequence 4, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FaastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 47; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.49;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
| | | | | | | | | |
DB 127 QLCTELQTT 135

RESULT 16

US-09-485-885-10
Sequence 10, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FaastSeq for Windows Version 3.0
SEQ ID NO 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 47; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.52;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
| | | | | | | | | |
DB 146 QLCTELQTT 154

RESULT 17

US-09-485-885-6
Sequence 6, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabazon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FaastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 47; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Db 127 QLCTELQTT 135

RESULT 18

US-09-485-885-14
; Sequence 14, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 47; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.69;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9

Db 146 QLCTELQTT 154

RESULT 19

US-09-854-133-425
; Sequence 425, Application US/09854133
; Patent No. 6759508
; GENERAL INFORMATION:
; APPLICANT: Lodes, Michael J.
; APPLICANT: Mohamath, Raodoh
; APPLICANT: Henderson, Robert A.
; APPLICANT: Benson, Darin R.
; APPLICANT: Secrist, Heather
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR
; FILE REFERENCE: 210121.475C10
; CURRENT APPLICATION NUMBER: US/09/854,133
; CURRENT FILING DATE: 2001-05-11
; NUMBER OF SEQ ID NOS: 735
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 425
; LENGTH: 4019
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-854-133-425

Query Match 72.3%; Score 34; DB 2; Length 4019;
Best Local Similarity 85.7%; Pred. No. 1.5e+03;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQ 7

Db 2730 QLCTEL 2736

RESULT 20
US-08-466-285-2
; Sequence 2, Application US/08466285

; Patent No. 5753233
; GENERAL INFORMATION:
; APPLICANT: Bleul, Conrad
; APPLICANT: Gissmann, Lutz
; APPLICANT: Muller, Martin
; TITLE OF INVENTION: Seroreactive Epitopes On Proteins Of
; TITLE OF INVENTION: Human Papillomavirus (HPV)18
; NUMBER OF SEQUENCES: 7
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; STREET: 1300 I Street, N.W., Suite 700
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION NUMBER: US/08/466,285
; FILING DATE: 06-JUN-1995
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/164,768
; FILING DATE: 10-DEC-1993
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/947,992
; FILING DATE: 21-SEP-1992
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/696,953
; FILING DATE: 08-MAY-1991
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: P 40 15 044.5
; FILING DATE: 10-MAY-1990
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: Manspeizer, David A.
; REGISTRATION NUMBER: 37,540
; REFERENCE/DOCKET NUMBER: 05552.1075-03000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (202)408-4000
; TELEFAX: (202)408-4400
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 32 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-466-285-2

Query Match 70.2%; Score 33; DB 1; Length 32;
Best Local Similarity 75.0%; Pred. No. 20;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9

Db 12 LCTELQTT 19

RESULT 21
US-08-164-768-2
; Sequence 2, Application US/08164768
; Patent No. 6322794
; GENERAL INFORMATION:
; APPLICANT: BLEUL, Conrad
; APPLICANT: GISSMANN, Lutz

APPLICANT: MULLER, Martin
TITLE OF INVENTION: SEROREACTIVE EPITOPES ON PROTEINS OF
TITLE OF INVENTION: HUMAN PAPILLOMA VIRUS (HPV) 18
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT &
ADDRESSEE: DUNNER, L.L.P.
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Forman, David S.
REGISTRATION NUMBER: 33,694
REFERENCE/DOCKET NUMBER: 05552.1075-02000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-164-768-2

Query Match 70.2%; Score 33; DB 2; Length 32;
Best Local Similarity 75.0%; Pred. No. 20;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
DB 12 LCTELNTS 19
RESULT 22
US-09-270-767-33527
Sequence 33527, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 33527
LENGTH: 141
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-33527

Query Match 70.2%; Score 33; DB 2; Length 141;
Best Local Similarity 85.7%; Pred. No. 85;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQ 7
DB 45 KLCTELQ 51

RESULT 23
US-09-270-767-48744
Sequence 48744, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 48744
LENGTH: 141
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-48744

Query Match 70.2%; Score 33; DB 2; Length 141;
Best Local Similarity 85.7%; Pred. No. 85;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCTELQ 7
DB 45 KLCTELQ 51

RESULT 24
US-08-247-904B-10
Sequence 10, Application US/08247904B
Patent No. 5981699
GENERAL INFORMATION:
APPLICANT: Rolfe, Mark
APPLICANT: Eckstein, Jens W.
APPLICANT: Draetta, Giulio
TITLE OF INVENTION: Human Ubiquitin Conjugating Enzyme
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley, Hoag & Eliot
STREET: One Post Office Square
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII(text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/247,904B
FILING DATE: 23-MAY-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: MIV-029.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 832-1000
TELEFAX: (617) 832-7000
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-247-904B-10

Query Match 70.2%; Score 33; DB 1; Length 158;
Best Local Similarity 75.0%; Pred. No. 95;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9

Db 17 LCTEIMTS 24

RESULT 25

US-08-767-942A-19
Sequence 19, Application US/08767942A

Patent No. 6068982

GENERAL INFORMATION:

APPLICANT: Rolfe, Mark

APPLICANT: Chiu, M. Isabel

APPLICANT: Berlin, Vivian

APPLICANT: Damaguez, Veronique

APPLICANT: Draetta, Giulio

APPLICANT: Guillaume, Cottarel

TITLE OF INVENTION: UBQUITIN CONJUGATING ENZYMES

NUMBER OF SEQUENCES: 45

CORRESPONDENCE ADDRESS:

ADDRESSEE: FOLEY, HONG & ELIOT LLP

STREET: One Post Office Square

CITY: Boston

STATE: MA

COUNTRY: USA

ZIP: 02109-2170

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/767,942A

FILING DATE: 17-DEC-1996

ATTORNEY/AGENT INFORMATION:

NAME: Vincent, Matthew P.

REGISTRATION NUMBER: 36,709

REFERENCE/DOCKET NUMBER: MIV-029.04

TELEPHONE: 617-832-1000

TELEFAX: 617-832-7000

INFORMATION FOR SEQ ID NO: 19:

SEQUENCE CHARACTERISTICS:

LENGTH: 158 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-767-942A-19

Query Match 70.2%; Score 33; DB 2; Length 158;
Best Local Similarity 75.0%; Pred. No. 95;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTEIMTS 9
Db 17 LCTEIMTS 24

RESULT 26

US-08-117-083-14

Sequence 14, Application US/08117083

Patent No. 5719054

GENERAL INFORMATION:

APPLICANT: Bournelli, Michael E.

APPLICANT: Ingile, Stephen C.

APPLICANT: Munro, Alan J.

TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human

NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:

ADDRESSEE: Walter H. Dreger

STREET: 4 Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: CA

COUNTRY: USA

ZIP: 94111

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/117,083

FILING DATE: 10-SEP-1993

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Dreger, Walter H.

REGISTRATION NUMBER: 24,190

REFERENCE/DOCKET NUMBER: A-58783

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-781-1989

TELEFAX: 415-398-3249

INFORMATION FOR SEQ ID NO: 14:

SEQUENCE CHARACTERISTICS:

LENGTH: 271 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

FEATURE: NAME/KEY: Protein

LOCATION: 1..271

OTHER INFORMATION: /note= "Xaa refers to stop codon in

US-08-117-083-14

Query Match 70.2%; Score 33; DB 1; Length 271;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTEIMTS 9
Db 18 LCTEIMTS 25

RESULT 27

US-09-485-885-21

Sequence 21, Application US/09485885

Patent No. 6342224

GENERAL INFORMATION:

APPLICANT: Bruck, Claudine

APPLICANT: Cabazon Silva, Teresa

APPLICANT: Delisse, Anne-Marie Eva Bernande

APPLICANT: Gerard, Catherine Marie Ghislaine

APPLICANT: Lombardo-Benchelkh, Angela

TITLE OF INVENTION: Vaccine

FILE REFERENCE: B45107

CURRENT APPLICATION NUMBER: US/09/485,885

PRIOR FILING DATE: 2000-02-18

PRIOR APPLICATION NUMBER: PCT/EP98/05285

PRIOR FILING DATE: 1998-08-17

PRIOR APPLICATION NUMBER: GB 9717953.5

PRIOR FILING DATE: 1997-08-22

NUMBER OF SEQ ID NOS: 23

SOFTWARE: FastSeq for Windows Version 3.0

SEQ ID NO 21

LENGTH: 278

TYPE: PRP

ORGANISM: Homo sapien

US-09-485-885-21

Query Match 70.2%; Score 33; DB 2; Length 278;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTEIMTS 9
Db 17 LCTEIMTS 24

```
Db          128 LCTELNTS 135

RESULT 28
US-09-485-885-23
; Sequence 23, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485, 885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP96/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 23
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-485-885-23

Query Match          70.2%; Score 33; DB 2; Length 383;
Best Local Similarity 75.0%; Pred. No. 2.3e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY          2 LCTELQTT 9
          |||||
          |||||

Db          128 LCTELNTS 135

RESULT 29
US-10-104-047-2475
; Sequence 2475, Application US/10104047
; Patent No. 6943241
; GENERAL INFORMATION:
; APPLICANT: HELIX RESEARCH INSTITUTE
; TITLE OF INVENTION: No. 6943241e1 full length cDNA
; FILE REFERENCE: H1-A0105
; CURRENT APPLICATION NUMBER: US/10/104, 047
; CURRENT FILING DATE: 2002-03-25
; PRIOR APPLICATION NUMBER:
; PRIOR FILING DATE:
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2475
; LENGTH: 1098
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-104-047-2475

Query Match          70.2%; Score 33; DB 2; Length 1098;
Best Local Similarity 85.7%; Pred. No. 6.4e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY          1 QLCSELO 7
          |||||
          |||||

Db          285 QLCSELO 291

RESULT 30
US-09-793-998-11
; Sequence 11, Application US/09793998
; Patent No. 6593119
; GENERAL INFORMATION:
; APPLICANT: KORCZAK, BOZENA

Db          175 QICAEQTT 182

Query Match          70.2%; Score 33; DB 2; Length 1104;
Best Local Similarity 62.5%; Pred. No. 6.4e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY          1 QLCSELO 8
          |||||
          |||||

Db          175 QICAEQTT 182

RESULT 31
US-08-787-547-101
; Sequence 101, Application US/08787547
; Patent No. 5783567
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Curley, Joanne M.
; APPLICANT: Langer, Robert S.
; TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY
; TITLE OF INVENTION: OF NUCLEIC ACID
; NUMBER OF SEQUENCES: 107
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Fish & Richardson, P.C.
; STREET: 225 Franklin Street
; CITY: Boston
; STATE: MA
; COUNTRY: US
; ZIP: 02110-2804
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: Windows95
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/787,547
; FILING DATE: 22-JAN-1997
; CLASSIFICATION: 514
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER:
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Fraser, Janis K.
; REGISTRATION NUMBER: 34,819
; REFERENCE/DOCKET NUMBER: 08191/003001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-542-5070
; TELEFAX: 617-542-8906
; TELEX: 200154
; INFORMATION FOR SEQ ID NO: 101:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 9 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-787-547-101

Query Match          68.1%; Score 32; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
```

Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTEL 6
| | | | |
Db 4 QCTEL 9

RESULT 32

US-09-601-729-274
; Sequence 274, Application US/09601729

; Patent No. 6683052

; GENERAL INFORMATION:

; APPLICANT: THIAM, KADER

; APPLICANT: AURIAULT, CLAUDE

; APPLICANT: GRAS-MASSÉ, HELENE

; APPLICANT: LOING, ESTELLE

; APPLICANT: VERWERDE, CLAUDE

; APPLICANT: GUILLET, JEAN GERARD

; TITLE OF INVENTION: LIPOPEPTIDES CONTAINING AN INTERFERON FRAGMENT AND USES

; FILE REFERENCE: USB-97-AU-IN

; CURRENT APPLICATION NUMBER: US/09/601,729

; PRIOR FILING DATE: 2000-11-20

; PRIOR APPLICATION NUMBER: PCT/FR99/00259

; PRIOR FILING DATE: 1999-02-05

; PRIOR APPLICATION NUMBER: 98 01439

; PRIOR FILING DATE: 1998-02-06

; NUMBER OF SEQ ID NOS: 281

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 274

; LENGTH: 9

; TYPE: PRT

; ORGANISM: Artificial Sequence

; FEATURE: Description of Artificial Sequence: Synthetic

; OTHER INFORMATION: peptide

US-09-601-729-274

Query Match 68.1%; Score 32; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTEL 6
| | | | |
Db 4 QCTEL 9

RESULT 33

US-09-248-796A-17743
; Sequence 17743, Application US/09248796A

; Patent No. 6747137

; GENERAL INFORMATION:

; APPLICANT: Keith weinstock et al

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN

; FILE REFERENCE: 107196.132

; CURRENT APPLICATION NUMBER: US/09/248,796A

; PRIOR FILING DATE: 1999-02-12

; PRIOR APPLICATION NUMBER: US 60/074,725

; PRIOR FILING DATE: 1998-02-13

; PRIOR APPLICATION NUMBER: US 60/096,409

; PRIOR FILING DATE: 1998-08-13

; NUMBER OF SEQ ID NOS: 28208

; SEQ ID NO 17743

; LENGTH: 203

; TYPE: PRT

; ORGANISM: Candida albicans

US-09-248-796A-17743

Query Match 68.1%; Score 32; DB 2; Length 203;
Best Local Similarity 87.5%; Pred. No. 1.8e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 QCTEL 8
| | | | |
Db 102 QCTEL 109

RESULT 34

US-09-543-681A-8111

; Sequence 8111, Application US/09543681A

; Patent No. 6605709

; GENERAL INFORMATION:

; APPLICANT: GARY BRETON

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS

; FILE REFERENCE: 2709.1002-001

; CURRENT APPLICATION NUMBER: US/09/543,681A

; PRIOR FILING DATE: 2000-04-05

; PRIOR APPLICATION NUMBER: US 60/128,706

; PRIOR FILING DATE: 1999-04-09

; NUMBER OF SEQ ID NOS: 8344

; SEQ ID NO 8111

; LENGTH: 211

; TYPE: PRT

; ORGANISM: Proteus mirabilis

US-09-543-681A-8111

Query Match 68.1%; Score 32; DB 2; Length 211;
Best Local Similarity 100.0%; Pred. No. 1.9e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTEL 6
| | | | |
Db 190 QCTEL 195

RESULT 35

US-09-634-137-32

; Sequence 32, Application US/09634137

; Patent No. 663265

; GENERAL INFORMATION:

; APPLICANT: Perrino, Fred W

; TITLE OF INVENTION: Mammalian Genes Encoding 3'-5' Exonuclease

; FILE REFERENCE: wak200/48001/4-018

; CURRENT APPLICATION NUMBER: US/09/634,137

; PRIOR FILING DATE: 2000-08-08

; PRIOR APPLICATION NUMBER: US 60/148,018

; PRIOR FILING DATE: 1999-08-09

; NUMBER OF SEQ ID NOS: 34

; SOFTWARE: PatentIn Ver. 2.1

; SEQ ID NO 32

; LENGTH: 236

; TYPE: PRT

; ORGANISM: Mus musculus

US-09-634-137-32

Query Match 68.1%; Score 32; DB 2; Length 236;
Best Local Similarity 100.0%; Pred. No. 2.1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 2 LCTEL 7
| | | | |
Db 127 LCTEL 132

RESULT 36

US-09-270-767-33015

; Sequence 33015, Application US/09270767

; Patent No. 6703491

; GENERAL INFORMATION:

; APPLICANT: Homburger et al.

; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster

; FILE REFERENCE: File Reference: 7326-094

; CURRENT APPLICATION NUMBER: US/09/270,767

; CURRENT FILING DATE: 1999-03-17

NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 33015
LENGTH: 251
TYPE: PRT
ORGANISM: Drosophila melanogaster
FEATURE:
OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-33015

Query Match 68.1%; Score 32; DB 2; Length 251;
Best Local Similarity 66.7%; Pred. No. 2.3e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 238 ELCTTLKTT 246

RESULT 37
US-09-543-681A-5423
Sequence 5423, Application US/09543681A
Patent No. 6605709
GENERAL INFORMATION:
APPLICANT: GARY BERTON
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
FILE REFERENCE: 2709.1002-001
CURRENT APPLICATION NUMBER: US/09/543, 681A
CURRENT FILING DATE: 2000-04-05
PRIOR APPLICATION NUMBER: US 60/128,706
PRIOR FILING DATE: 1999-04-09
NUMBER OF SEQ ID NOS: 8344
SEQ ID NO 5423
LENGTH: 469
TYPE: PRT
ORGANISM: Proteus mirabilis
US-09-543-681A-5423

Query Match 68.1%; Score 32; DB 2; Length 469;
Best Local Similarity 75.0%; Pred. No. 4.2e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 8
Db 288 ELCTTLKTT 295

RESULT 38
US-09-121-964-9
Sequence 9, Application US/09121964
Patent No. 6124447
GENERAL INFORMATION:
APPLICANT: Natori, Shunji
TITLE OF INVENTION: NOVEL ENZYME CATALYZING DEPHOSPHORYLATION
FILE REFERENCE: 32290-144753
CURRENT APPLICATION NUMBER: US/09/121,964
CURRENT FILING DATE: 1998-07-24
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 9
LENGTH: 771
TYPE: PRT
ORGANISM: C. elegans
US-09-121-964-9

Query Match 68.1%; Score 32; DB 2; Length 771;
Best Local Similarity 77.8%; Pred. No. 6.8e+02;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 421 QLTSLQTT 429

RESULT 39
US-08-159-339A-320

Sequence 320, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esteban
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993

ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300

TELEX:
INFORMATION FOR SEQ ID NO: 320:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-320

Query Match 66.0%; Score 31; DB 2; Length 9;
Best Local Similarity 71.4%; Pred. No. 4.6e+05;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQ 7
Db 3 QMCTELK 9

RESULT 40
US-08-271-539-3
Sequence 3, Application US/08271539
Patent No. 6358509

GENERAL INFORMATION:
APPLICANT: Ramathanan, Lata
APPLICANT: Seelig, Gail F.
APPLICANT: Trotta, Paul P.
TITLE OF INVENTION: Antibody Antagonists of Human Interleukin-4
FILE REFERENCE: JB0059KQ US

CURRENT APPLICATION NUMBER: US/08/271,539
 CURRENT FILING DATE: 1994-07-07
 PRIOR APPLICATION NUMBER: US 07/453,570
 PRIOR FILING DATE: 1989-12-20
 PRIOR APPLICATION NUMBER: PCT/US90/07289
 PRIOR FILING DATE: 1990-12-18
 PRIOR APPLICATION NUMBER: US 07/859,689
 PRIOR FILING DATE: 1992-06-11
 NUMBER OF SEQ ID NOS: 46
 SOFTWARE: PatentIn Ver. 2.1
 SEQ ID NO: 3
 LENGTH: 14
 TYPE: PRT
 ORGANISM: Artificial Sequence
 FEATURE:
 OTHER INFORMATION: Description of Artificial Sequence: synthetic
 US-08-271-539-3

Query Match 66.0%; Score 31; DB 2; Length 14;
 Best Local Similarity 75.0%; Pred. No. 20;
 Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
 DB 6 LCTELTVT 13

RESULT 41
 US-09-513-999C-6885
 Sequence 6885, Application US/09513999C
 Patent No. 6783961
 GENERAL INFORMATION:
 APPLICANT: Dumas Milne Edwards, J.B.
 APPLICANT: Duclercq, A.
 APPLICANT: Giordano, J.Y.
 TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
 Patent No. 6783961
 FILE REFERENCE: 59 US2, REG
 CURRENT APPLICATION NUMBER: US/09/513,999C
 CURRENT FILING DATE: 2000-02-24
 PRIOR APPLICATION NUMBER: US 60/122,487
 PRIOR FILING DATE: 1999-02-26
 NUMBER OF SEQ ID NOS: 36681
 SOFTWARE: Patent.pm
 SEQ ID NO 6885
 LENGTH: 56
 TYPE: PRT
 ORGANISM: Homo sapiens
 US-09-513-999C-6885

Query Match 66.0%; Score 31; DB 2; Length 56;
 Best Local Similarity 71.4%; Pred. No. 78;
 Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 OLCTELO 7
 DB 33 QVCTOLO 39

RESULT 42
 US-08-234-812-3
 Sequence 3, Application US/08234812
 Patent No. 5557535
 GENERAL INFORMATION:
 APPLICANT: Srinivasan, Subhashini
 APPLICANT: Sudarshanam, Padmanaban
 TITLE OF INVENTION: METHOD AND SYSTEM FOR PROTEIN MODELING
 NUMBER OF SEQUENCES: 3
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Seed and Berry
 STREET: 6300 Columbia Center, 701 Fifth Avenue
 CITY: Seattle

STATE: Washington
 COUNTRY: US
 ZIP: 98104-7092
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/234,812
 FILING DATE: 28-APR-1994
 CLASSIFICATION: 395
 ATTORNEY/AGENT INFORMATION:
 NAME: Pirio, Maurice J.
 REGISTRATION NUMBER: 33,273
 REFERENCE/DOCKET NUMBER: 480052.408C1
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (206) 622-4900
 TELEFAX: (206) 682-6031
 TELEX: 3723836 SEDBANBRY
 INFORMATION FOR SEQ ID NO: 3:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 108 amino acids
 TYPE: amino acid
 STRANDEDNESS: single
 TOPOLOGY: linear
 US-08-234-812-3

Query Match 66.0%; Score 31; DB 1; Length 108;
 Best Local Similarity 75.0%; Pred. No. 1.5e+02;
 Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
 DB 17 LCTELTVT 24

RESULT 43
 US-08-663-809-3
 Sequence 3, Application US/08663809
 Patent No. 5884230
 GENERAL INFORMATION:
 APPLICANT: Srinivasan, Subhashini
 APPLICANT: Sudarshanam, Padmanaban
 TITLE OF INVENTION: METHOD AND SYSTEM FOR PROTEIN MODELING
 NUMBER OF SEQUENCES: 3
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Seed and Berry
 STREET: 6300 Columbia Center, 701 Fifth Avenue
 CITY: Seattle
 STATE: Washington
 COUNTRY: US
 ZIP: 98104-7092
 COMPUTER READABLE FORM:
 MEDIUM TYPE: Floppy disk
 COMPUTER: IBM PC compatible
 OPERATING SYSTEM: PC-DOS/MS-DOS
 SOFTWARE: PatentIn Release #1.0, Version #1.25
 CURRENT APPLICATION DATA:
 APPLICATION NUMBER: US/08/663,809
 FILING DATE: 14-JUN-1996
 CLASSIFICATION: 364
 ATTORNEY/AGENT INFORMATION:
 NAME: Pirio, Maurice J.
 REGISTRATION NUMBER: 33,273
 REFERENCE/DOCKET NUMBER: 480052.408C2
 TELECOMMUNICATION INFORMATION:
 TELEPHONE: (206) 622-4900
 TELEFAX: (206) 682-6031
 INFORMATION FOR SEQ ID NO: 3:
 SEQUENCE CHARACTERISTICS:
 LENGTH: 108 amino acids
 TYPE: amino acid

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; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-663-809-3

Query Match      66.0%; Score 31; DB 1; Length 108;
Best Local Similarity 75.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY      2 LCTELQTT 9
        |||||
Db      17 LCTELTFT 24

RESULT 44
US-08-049-503-1
; Sequence 1, Application US/08049503
; Patent No. 5494662
; GENERAL INFORMATION:
; APPLICANT: UENO, KOHJI
; APPLICANT: KATAYAMA, TERUOKI
; APPLICANT: MIYAMOTO, TSUMORU
; TITLE OF INVENTION: STIMULATOR FOR BONE FORMATION
; NUMBER OF SEQUENCES: 12
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: STEVENS, DAVIS, MILLER & MOSHER, LLP
; STREET: 515 N. WASHINGTON ST.
; CITY: ALEXANDRIA
; STATE: VA
; COUNTRY: USA
; ZIP: 22314
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/049,503
; FILING DATE: 21-APR-1993
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: POULOS III, JAMES A.
; REGISTRATION NUMBER: 31,714
; REFERENCE/DOCKET NUMBER: TPP29045
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 703-549-7200
; TELEFAX: 703-528-5313
; INFORMATION FOR SEQ ID NO: 1:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 129 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
; ANTI-SENSE: NO
; FRAGMENT TYPE: N-terminal
; ORIGINAL SOURCE:
; ORGANISM: Homo sapiens
US-08-049-503-1

Query Match      66.0%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY      2 LCTELQTT 9
        |||||
Db      23 LCTELTFT 30

RESULT 45
US-08-225-224-2
; Sequence 2, Application US/08225224
; Patent No. 5635599
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```

; GENERAL INFORMATION:
; APPLICANT: PASTAN, Ira
; APPLICANT: KREITMAN, Robert J.
; TITLE OF INVENTION: CIRCULARLY PERMUTATED LIGANDS AND
; TITLE OF INVENTION: CIRCULARLY PERMUTED FUSION PROTEINS
; NUMBER OF SEQUENCES: 57
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Townsend and Townsend Kourie and Crew
; STREET: Stewart Street Tower, One Market Plaza
; CITY: San Francisco
; STATE: California
; COUNTRY: US
; ZIP: 94105-1493
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/225,224
; FILING DATE: 8-APR-1994
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: Weber, Ellen L.
; REGISTRATION NUMBER: 32,762
; REFERENCE/DOCKET NUMBER: 15280-193
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 543-9600
; TELEFAX: (415) 543-5043
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 129 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..129
; OTHER INFORMATION: /label= IL4
US-08-225-224-2

Query Match      66.0%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY      2 LCTELQTT 9
        |||||
Db      23 LCTELTFT 30

RESULT 46
US-08-470-299-21
; Sequence 21, Application US/08470299
; Patent No. 5783181
; GENERAL INFORMATION:
; APPLICANT: Browne, Michael J.
; APPLICANT: Murphy, Kay B.
; APPLICANT: Chapman, Conrad G.
; APPLICANT: Clinkenbeard, Helen B.
; APPLICANT: Young, Peter R.
; APPLICANT: Shatzman, Allan R.
; TITLE OF INVENTION: No. 5783181el Compounds
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: SmithKline Beecham Corporation
; STREET: 709 Swedeland Road, P.O. Box 1539
; CITY: King of Prussia
; STATE: Pennsylvania
; COUNTRY: USA
; ZIP: 19406
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,299
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
US-08-470-299-21

Query Match 66.0%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||
DB 23 LCTELVT 30

RESULT 47
US-08-874-697-1
Sequence 1, Application US/08874697
Patent No. 5986059
GENERAL INFORMATION:
APPLICANT: Shanefelt, Armen; Greve, Jeffrey; Gandel, Robert
TITLE OF INVENTION: T-cell Selective Interleukin-4 Agonists
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bayer Corporation, Pharmaceutical Division
STREET: 400 Morgan Lane
CITY: West Haven
STATE: CT
COUNTRY: United States of America
ZIP: 06516-4175
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS V. 6.30
SOFTWARE: word for windows 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/874,697
FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/019,748
FILING DATE: 14-JUN-1996
PRIOR APPLICATION DATA: 60/036,746
FILING DATE: 27-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Huw R. Jones
REGISTRATION NUMBER: 33,916
REFERENCE/DOCKET NUMBER: WH5013
TELECOMMUNICATION INFORMATION:
TELEPHONE: (203) 812-2317
TELEFAX: (203) 812-5492
INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 129
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
DESCRIPTION: human Interleukin-4 protein
HYPOTHETICAL: no
ANTI-SENSE: no
US-08-874-697-1

Query Match 66.0%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||
DB 23 LCTELVT 30

RESULT 48
US-08-722-258-2
Sequence 2, Application US/08722258
Patent No. 6011002
GENERAL INFORMATION:
APPLICANT: Pastan, Ira
APPLICANT: Kreitman, Robert J.
APPLICANT: Puri, Raj K.
TITLE OF INVENTION: Circularly Permuted Ligands and
TITLE OF INVENTION: Circularly Permuted Chimeric Molecules
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/722,258
FILING DATE: 08-JAN-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/04468
FILING DATE: 06-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/225,224
FILING DATE: 08-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 015280-193100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..129
OTHER INFORMATION: /note="Interleukin 4 (IL4)"
US-08-722-258-2

Query Match 66.0%; Score 31; DB 2; Length 129;
 Best Local Similarity 75.0%; Pred. No. 1.8e+02;
 Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
 |||||
 Db 23 LCTELTVT 30

RESULT 49
 US-08-897-020-1
 ; Sequence 1, Application US/08897020
 ; Patent No. 6028176
 ; GENERAL INFORMATION:
 ; APPLICANT: Sharnfeldt, Armen; Greve, Jeffrey; Rocznik, Steven
 ; TITLE OF INVENTION: High-affinity Interleukin-4 Molecules
 ; NUMBER OF SEQUENCES: 9
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: Bayer Corporation, Pharmaceutical Division
 ; STREET: 400 Morgan Lane
 ; CITY: West Haven
 ; STATE: CT
 ; COUNTRY: United States of America
 ; ZIP: 06516-4175
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS v. 6.30
 ; SOFTWARE: word for windows 6.0
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/897,020
 ; FILING DATE:
 ; CLASSIFICATION: 435
 ; PRIOR APPLICATION DATA:
 ; APPLICATION NUMBER: P-91,242
 ; FILING DATE: 19-JUL-1996
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: Huw R. Jones
 ; REGISTRATION NUMBER: 33, 916
 ; REFERENCE/DOCKET NUMBER: WH5020
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: (203) 812-2317
 ; TELEFAX: (203) 812-5492
 ; INFORMATION FOR SEQ ID NO: 1:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 129
 ; TYPE: amino acid
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 ; MOLECULE TYPE: protein
 ; DESCRIPTION: human Interleukin-4 protein
 ; HYPOTHEICAL: no
 ; ANTI-SENSE: no
 ; US-08-897-020-1

Query Match 66.0%; Score 31; DB 2; Length 129;
 Best Local Similarity 75.0%; Pred. No. 1.8e+02;
 Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
 |||||
 Db 23 LCTELTVT 30

RESULT 50
 US-08-765-012A-16
 ; Sequence 16, Application US/08755012A
 ; Patent No. 6130318
 ; GENERAL INFORMATION:
 ; APPLICANT: Wild, Hanno; Hanko, Rudolf; Dorschug, Michael;
 ; APPLICANT: Horlein, Hans-Dietrich; Beunink, Jurgen;
 ; APPLICANT: Apeler, Heiner; Wehlmann, Hermann; and Sebald,
 ; APPLICANT: Walter

;; TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS
 ;; TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
 ;; TITLE OF INVENTION: HUMAN INTERLEUKIN 4
 ;; NUMBER OF SEQUENCES: 20
 ;; CORRESPONDENCE ADDRESS:
 ;; ADDRESSEE: Sprung Kramer Schaefer & Briscoe
 ;; STREET: 660 White Plains Road
 ;; CITY: Tarrytown
 ;; STATE: New York
 ;; COUNTRY: USA
 ;; ZIP: 10591-5144
 ;; COMPUTER READABLE FORM:
 ;; MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
 ;; COMPUTER: Apple Macintosh
 ;; OPERATING SYSTEM: System 7.5
 ;; SOFTWARE: WordPerfect 3.5
 ;; CURRENT APPLICATION DATA:
 ;; APPLICATION NUMBER: US/08/765,012A
 ;; FILING DATE: 19-DEC-1996
 ;; CLASSIFICATION: 435
 ;; PRIOR APPLICATION DATA:
 ;; APPLICATION NUMBER: PCT/EP95/02358
 ;; FILING DATE: 19-JUN-1995
 ;; PRIOR APPLICATION DATA:
 ;; APPLICATION NUMBER: DE 44 23 131
 ;; FILING DATE: 01-JUL-1994
 ;; ATTORNEY/AGENT INFORMATION:
 ;; NAME: Kute G. Briscoe
 ;; REGISTRATION NUMBER: 33,141
 ;; REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
 ;; TELECOMMUNICATION INFORMATION:
 ;; TELEPHONE: (914) 332-1700
 ;; TELEFAX: (914) 332-1844
 ;; INFORMATION FOR SEQ ID NO: 16:
 ;; SEQUENCE CHARACTERISTICS:
 ;; LENGTH: 129 amino acids
 ;; TYPE: amino acid
 ;; TOPOLOGY: linear
 ; US-08-765-012A-16

Query Match 66.0%; Score 31; DB 2; Length 129;
 Best Local Similarity 75.0%; Pred. No. 1.8e+02;
 Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
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 Db 23 LCTELTVT 30

Search completed: May 5, 2006, 06:24:13
 Job time : 24.9 secs

GenCore version 5.1.7
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OM protein - protein search, using SW model

Run on: May 5, 2006, 08:39:55 ; Search time 56.3 Seconds
(without alignments)
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Title: US-08-170-344-3
Perfect score: 47
Sequence: 1 QLTCLQRT 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

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6: /cgn2_6/ptodata/1/pubppaa/US11_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	47	100.0	15	4	US-10-476-570-20
2	47	100.0	15	4	US-10-476-570-21
3	47	100.0	21	4	US-10-476-570-8
4	47	100.0	25	6	US-11-021-949-1
5	47	100.0	30	4	US-10-476-570-53
6	47	100.0	30	5	US-10-858-384-4
7	47	100.0	32	4	US-10-476-570-9
8	47	100.0	33	4	US-10-476-570-19
9	47	100.0	151	5	US-10-177-390-6
10	47	100.0	151	5	US-10-484-063-27
11	47	100.0	151	5	US-10-484-063-20
12	47	100.0	158	5	US-10-858-384-2
13	47	100.0	158	5	US-10-367-057-16
14	47	100.0	158	6	US-11-021-949-13
15	47	100.0	171	4	US-10-472-724-2
16	47	100.0	243	6	US-11-072-288-1
17	47	100.0	266	3	US-09-367-309A-1
18	47	100.0	273	4	US-10-000-903-4
19	47	100.0	273	5	US-10-899-771-1
20	47	100.0	292	4	US-10-000-903-10
21	47	100.0	292	5	US-10-899-771-10
22	47	100.0	371	4	US-10-000-903-6
23	47	100.0	371	5	US-10-899-771-6
24	47	100.0	380	4	US-10-000-903-14
25	47	100.0	390	5	US-10-899-771-14
26	47	100.0	536	4	US-10-367-095-10
27	47	100.0	536	4	US-10-368-046-10

28	47	100.0	536	4	US-10-367-367-10	Sequence 10, Appl
29	47	100.0	536	5	US-10-918-337-10	Sequence 10, Appl
30	37	78.7	48	3	US-09-925-299-1386	Sequence 1386, Ap
31	37	78.7	48	3	US-09-925-299-1386	Sequence 1386, Ap
32	37	78.7	587	4	US-10-437-963-115784	Sequence 115784,
33	37	78.7	853	4	US-10-437-963-122232	Sequence 122232,
34	36	76.6	55	4	US-10-424-599-191823	Sequence 191823,
35	36	76.6	470	4	US-10-369-493-1712	Sequence 1712, Ap
36	35	74.5	158	4	US-10-282-122A-70249	Sequence 70249, A
37	35	74.5	158	5	US-10-857-625-567	Sequence 567, App
38	35	74.5	239	5	US-10-820-155-14	Sequence 14, Appl
39	35	74.5	239	5	US-10-820-155-23	Sequence 23, Appl
40	35	74.5	239	5	US-10-820-155-30	Sequence 30, Appl
41	35	74.5	239	5	US-10-820-155-39	Sequence 39, Appl
42	35	74.5	239	5	US-10-820-155-85	Sequence 85, Appl
43	35	74.5	239	5	US-10-820-155-94	Sequence 94, Appl
44	35	74.5	239	5	US-10-820-155-97	Sequence 97, Appl
45	34	72.3	104	4	US-10-424-599-246277	Sequence 246277,
46	34	72.3	120	4	US-10-767-701-37218	Sequence 37218, A
47	34	72.3	202	4	US-10-127-816-9	Sequence 9, Appl1
48	34	72.3	202	4	US-10-127-816-11	Sequence 11, Appl
49	34	72.3	202	4	US-10-142-717-12	Sequence 12, Appl
50	34	72.3	202	4	US-10-420-034A-60	Sequence 60, Appl
51	34	72.3	202	4	US-10-420-034A-62	Sequence 62, Appl
52	34	72.3	202	4	US-10-691-923-8	Sequence 8, Appl1
53	34	72.3	202	4	US-10-691-923-10	Sequence 10, Appl
54	34	72.3	202	5	US-10-914-772-8	Sequence 8, Appl1
55	34	72.3	202	5	US-10-914-772-10	Sequence 10, Appl
56	34	72.3	202	5	US-10-425-114-71026	Sequence 71026, A
57	34	72.3	342	3	US-09-886-055-207	Sequence 207, App
58	34	72.3	342	3	US-09-804-291-207	Sequence 207, App
59	34	72.3	342	5	US-10-819-316-207	Sequence 207, App
60	34	72.3	342	5	US-10-425-115-36747	Sequence 36747, A
61	34	72.3	378	4	US-10-425-114-63762	Sequence 63762, A
62	34	72.3	613	4	US-10-425-115-343502	Sequence 343502,
63	34	72.3	613	4	US-10-424-599-226123	Sequence 226123,
64	34	72.3	806	5	US-10-732-923-8378	Sequence 8378, Ap
65	34	72.3	4019	3	US-09-738-927-425	Sequence 425, App
66	34	72.3	4019	3	US-09-854-133-425	Sequence 425, App
67	34	72.3	4019	4	US-10-144-649A-425	Sequence 425, App
68	33	70.2	189	4	US-10-424-599-192924	Sequence 192924,
69	33	70.2	25	6	US-11-021-949-2	Sequence 2, Appl1
70	33	70.2	25	6	US-11-021-949-57	Sequence 57, Appl
71	33	70.2	42	5	US-10-751-845-152	Sequence 152, App
72	33	70.2	53	4	US-10-437-963-184267	Sequence 184267,
73	33	70.2	81	4	US-10-424-599-222742	Sequence 222742,
74	33	70.2	82	4	US-10-437-963-175677	Sequence 175677,
75	33	70.2	119	5	US-10-751-845-159	Sequence 159, App
76	33	70.2	137	4	US-10-751-845-159	Sequence 159, App
77	33	70.2	158	5	US-10-800-022-27	Sequence 27, Appl
78	33	70.2	158	6	US-11-021-949-28	Sequence 28, Appl
79	33	70.2	158	6	US-11-021-949-29	Sequence 29, Appl
80	33	70.2	172	4	US-10-472-724-6	Sequence 6, Appl1
81	33	70.2	179	4	US-10-424-599-147599	Sequence 147599,
82	33	70.2	236	5	US-10-751-845-157	Sequence 157, App
83	33	70.2	237	5	US-10-751-845-158	Sequence 158, App
84	33	70.2	252	4	US-10-767-701-43809	Sequence 43809, A
85	33	70.2	256	4	US-10-424-599-147707	Sequence 147707,
86	33	70.2	258	4	US-10-739-930-6593	Sequence 6593, Ap
87	33	70.2	261	5	US-10-751-845-160	Sequence 160, App
88	33	70.2	278	4	US-10-000-903-21	Sequence 21, Appl
89	33	70.2	278	5	US-10-899-771-21	Sequence 21, Appl
90	33	70.2	312	4	US-10-424-599-147488	Sequence 147488,
91	33	70.2	312	4	US-10-424-599-147707	Sequence 147707,
92	33	70.2	337	4	US-10-425-114-66915	Sequence 66915, A
93	33	70.2	342	4	US-10-425-114-47110	Sequence 47110, A
94	33	70.2	383	4	US-10-000-903-23	Sequence 23, Appl
95	33	70.2	383	5	US-10-899-771-23	Sequence 23, Appl
96	33	70.2	393	5	US-10-954-778-41	Sequence 41, Appl
97	33	70.2	562	4	US-10-424-599-251264	Sequence 251264,
98	33	70.2	662	6	US-11-097-143-12708	Sequence 12708, A
99	33	70.2	759	4	US-10-408-765A-748	Sequence 748, App
100	33	70.2	863	4	US-10-437-963-194291	Sequence 194291,

101	33	70.2	1098	4	US-10-104-047-2475	Sequence 2475, App	174	31	66.0	129	4	US-10-658-834A-567	Sequence 567, App
102	33	70.2	1104	3	US-09-793-998-11	Sequence 11, Appl	175	31	66.0	129	4	US-10-773-939-11	Sequence 11, Appl
103	33	70.2	1104	3	US-10-423-893-11	Sequence 11, Appl	176	31	66.0	129	4	US-10-773-939-11	Sequence 11, Appl
104	33	68.1	1104	9	US-09-909-460-101	Sequence 101, App	177	31	66.0	129	4	US-10-773-939-11	Sequence 11, Appl
105	32	68.1	9	3	US-09-872-836-101	Sequence 101, App	178	31	66.0	129	5	US-10-866-540-11	Sequence 11, Appl
106	32	68.1	9	4	US-10-133-210-278	Sequence 278, App	179	31	66.0	129	5	US-10-866-540-11	Sequence 11, Appl
107	32	68.1	9	4	US-10-777-053-546	Sequence 546, App	180	31	66.0	129	5	US-10-872-198-118	Sequence 118, App
108	32	68.1	9	4	US-10-837-217-546	Sequence 546, App	181	31	66.0	129	5	US-10-873-332-63	Sequence 63, Appl
109	32	68.1	9	5	US-10-738-970-101	Sequence 101, App	182	31	66.0	129	5	US-10-605-288-11	Sequence 11, Appl
110	32	68.1	9	5	US-10-484-063-1	Sequence 1, Appl1	183	31	66.0	129	5	US-10-866-580-11	Sequence 11, Appl
111	32	68.1	9	5	US-10-751-845-55	Sequence 55, Appl	184	31	66.0	129	5	US-10-773-930-11	Sequence 11, Appl
112	32	68.1	12	6	US-11-021-949-7	Sequence 7, Appl1	185	31	66.0	129	6	US-11-021-951-118	Sequence 118, App
113	32	68.1	20	5	US-10-751-845-64	Sequence 64, Appl1	186	31	66.0	129	6	US-11-071-098-11	Sequence 11, Appl
114	32	68.1	60	4	US-10-424-599-202842	Sequence 202842, App	187	31	66.0	129	6	US-11-070-993-11	Sequence 11, Appl
115	32	68.1	76	3	US-09-867-550-1312	Sequence 1312, App	188	31	66.0	129	6	US-11-112-701-16	Sequence 16, Appl
116	32	68.1	82	4	US-10-424-599-179241	Sequence 179241, App	189	31	66.0	130	4	US-10-033-355A-4	Sequence 4, Appl1
117	32	68.1	88	4	US-10-425-115-362556	Sequence 362556, App	190	31	66.0	130	5	US-10-820-559-9	Sequence 9, Appl1
118	32	68.1	114	6	US-11-097-143-36090	Sequence 36090, A	191	31	66.0	130	5	US-10-820-559-11	Sequence 11, Appl
119	32	68.1	117	5	US-10-751-845-126	Sequence 126, App	192	31	66.0	130	5	US-10-820-559-12	Sequence 12, Appl
120	32	68.1	212	4	US-10-363-616-453	Sequence 453, App	193	31	66.0	130	5	US-10-820-559-13	Sequence 13, Appl
121	32	68.1	227	3	US-09-925-302-604	Sequence 604, App	194	31	66.0	130	5	US-10-820-559-14	Sequence 14, Appl
122	32	68.1	227	3	US-09-925-302-604	Sequence 604, App	195	31	66.0	130	5	US-10-820-559-15	Sequence 15, Appl
123	32	68.1	265	4	US-10-354-437-30	Sequence 30, Appl	196	31	66.0	130	5	US-10-820-559-16	Sequence 16, Appl
124	32	68.1	341	5	US-10-450-763-54568	Sequence 54568, A	197	31	66.0	139	4	US-10-425-115-229668	Sequence 229668, App
125	32	68.1	351	4	US-10-369-493-3387	Sequence 3387, App	198	31	66.0	150	4	US-10-233-902-25	Sequence 25, Appl
126	32	68.1	358	4	US-10-425-115-288633	Sequence 288633, App	199	31	66.0	150	6	US-11-041-636-25	Sequence 25, Appl
127	32	68.1	398	5	US-10-934-778-39	Sequence 39, Appl	200	31	66.0	153	3	US-09-947-770-13	Sequence 13, Appl
128	32	68.1	418	4	US-10-282-122A-69665	Sequence 69665, A	201	31	66.0	153	3	US-09-923-246-112	Sequence 112, App
129	32	68.1	452	5	US-10-454-763-37868	Sequence 37868, A	202	31	66.0	153	4	US-10-295-723-112	Sequence 112, App
130	32	68.1	555	4	US-10-094-749-2279	Sequence 2279, App	203	31	66.0	153	4	US-10-282-622-8	Sequence 8, Appl1
131	32	68.1	560	4	US-10-363-829-322	Sequence 322, App	204	31	66.0	153	4	US-10-351-157-179	Sequence 179, App
132	32	68.1	574	4	US-10-437-963-102998	Sequence 102998, App	205	31	66.0	153	4	US-10-352-554-164	Sequence 164, App
133	32	68.1	718	4	US-10-087-192-1680	Sequence 1680, App	206	31	66.0	153	4	US-10-456-780-8	Sequence 8, Appl1
134	32	68.1	751	4	US-10-369-493-3357	Sequence 3357, App	207	31	66.0	153	4	US-10-659-684-112	Sequence 112, App
135	32	68.1	1354	4	US-10-389-566-1269	Sequence 1269, App	208	31	66.0	153	4	US-10-683-516-1	Sequence 1, Appl1
136	32	68.1	1354	4	US-10-437-963-176978	Sequence 176978, App	209	31	66.0	153	5	US-10-688-845-6	Sequence 6, Appl1
137	31	66.0	452	4	US-10-425-115-383541	Sequence 283541, App	210	31	66.0	153	5	US-10-688-845-6	Sequence 6, Appl1
138	31	66.0	85	4	US-10-437-963-156344	Sequence 156344, App	211	31	66.0	153	5	US-10-787-442-112	Sequence 112, App
139	31	66.0	95	4	US-10-425-115-212369	Sequence 212369, App	212	31	66.0	153	5	US-10-775-204-2178	Sequence 2178, App
140	31	66.0	107	4	US-10-437-963-165626	Sequence 165626, App	213	31	66.0	153	5	US-10-775-204-2178	Sequence 2178, App
141	31	66.0	109	5	US-10-450-763-46825	Sequence 46825, A	214	31	66.0	158	6	US-11-021-949-301	Sequence 301, Appl
142	31	66.0	128	4	US-10-150-874-2	Sequence 2, Appl1	215	31	66.0	160	4	US-10-424-559-171507	Sequence 171507, App
143	31	66.0	129	3	US-09-792-793A-25	Sequence 25, Appl	216	31	66.0	160	4	US-10-424-559-171507	Sequence 171507, App
144	31	66.0	129	3	US-09-969-748C-27	Sequence 27, Appl	217	31	66.0	160	4	US-10-424-559-171507	Sequence 171507, App
145	31	66.0	129	3	US-10-040-586-1	Sequence 1, Appl1	218	31	66.0	214	3	US-09-969-836-1	Sequence 1, Appl1
146	31	66.0	129	4	US-10-050-227-21	Sequence 21, Appl	219	31	66.0	221	6	US-11-097-143-29653	Sequence 29653, App
147	31	66.0	129	4	US-10-400-377-11	Sequence 11, Appl	220	31	66.0	220	5	US-10-181-5858-114	Sequence 114, App
148	31	66.0	129	4	US-10-400-708-11	Sequence 11, Appl	221	31	66.0	230	5	US-10-181-5858-118	Sequence 118, App
149	31	66.0	129	4	US-10-298-148-11	Sequence 11, Appl	222	31	66.0	231	5	US-10-732-923-2569	Sequence 2569, App
150	31	66.0	129	4	US-10-375-209A-25	Sequence 25, Appl	223	31	66.0	231	5	US-10-181-5858-107	Sequence 107, App
151	31	66.0	129	4	US-10-280-671-1	Sequence 1, Appl1	224	31	66.0	231	5	US-10-181-5858-111	Sequence 111, App
152	31	66.0	129	4	US-10-658-834A-207	Sequence 207, App	225	31	66.0	244	5	US-10-820-155-106	Sequence 106, App
153	31	66.0	129	4	US-10-658-834A-546	Sequence 546, App	226	31	66.0	245	4	US-10-282-122A-74397	Sequence 74397, A
154	31	66.0	129	4	US-10-658-834A-547	Sequence 547, App	227	31	66.0	245	5	US-10-181-5858-112	Sequence 112, App
155	31	66.0	129	4	US-10-658-834A-548	Sequence 548, App	228	31	66.0	245	5	US-10-181-5858-116	Sequence 116, App
156	31	66.0	129	4	US-10-658-834A-548	Sequence 548, App	229	31	66.0	246	5	US-10-181-5858-105	Sequence 105, App
157	31	66.0	129	4	US-10-658-834A-550	Sequence 550, App	230	31	66.0	246	5	US-10-181-5858-109	Sequence 109, App
158	31	66.0	129	4	US-10-658-834A-551	Sequence 551, App	231	31	66.0	244	4	US-10-006-867-62	Sequence 62, Appl
159	31	66.0	129	4	US-10-658-834A-552	Sequence 552, App	232	31	66.0	244	4	US-10-052-586-236	Sequence 236, App
160	31	66.0	129	4	US-10-658-834A-553	Sequence 553, App	233	31	66.0	244	4	US-10-063-547-62	Sequence 62, Appl
161	31	66.0	129	4	US-10-658-834A-554	Sequence 554, App	234	31	66.0	244	4	US-10-063-551-62	Sequence 62, Appl
162	31	66.0	129	4	US-10-658-834A-555	Sequence 555, App	235	31	66.0	244	4	US-10-174-590-236	Sequence 236, App
163	31	66.0	129	4	US-10-658-834A-556	Sequence 556, App	236	31	66.0	244	4	US-10-176-758-236	Sequence 236, App
164	31	66.0	129	4	US-10-658-834A-557	Sequence 557, App	237	31	66.0	244	4	US-10-175-737-236	Sequence 236, App
165	31	66.0	129	4	US-10-658-834A-558	Sequence 558, App	238	31	66.0	244	4	US-10-063-616-62	Sequence 62, Appl
166	31	66.0	129	4	US-10-658-834A-559	Sequence 559, App	239	31	66.0	244	4	US-10-174-581-236	Sequence 236, App
167	31	66.0	129	4	US-10-658-834A-560	Sequence 560, App	240	31	66.0	244	4	US-10-176-483-236	Sequence 236, App
168	31	66.0	129	4	US-10-658-834A-561	Sequence 561, App	241	31	66.0	244	4	US-10-176-914-236	Sequence 236, App
169	31	66.0	129	4	US-10-658-834A-562	Sequence 562, App	242	31	66.0	244	4	US-10-176-914-236	Sequence 236, App
170	31	66.0	129	4	US-10-658-834A-563	Sequence 563, App	243	31	66.0	244	4	US-10-176-915-236	Sequence 236, App
171	31	66.0	129	4	US-10-658-834A-564	Sequence 564, App	244	31	66.0	244	4	US-10-063-569-62	Sequence 62, Appl
172	31	66.0	129	4	US-10-658-834A-565	Sequence 565, App	245	31	66.0	244	4	US-10-063-513-62	Sequence 62, Appl
173	31	66.0	129	4	US-10-658-834A-566	Sequence 566, App	246	31	66.0	244	4	US-10-063-515-62	Sequence 62, Appl

393	31	66.0	284	4	US-10-196-745-236	Sequence 236, App	466	31	66.0	284	4	US-10-184-621-236	Sequence 236, App
394	31	66.0	284	4	US-10-196-762-236	Sequence 236, App	467	31	66.0	284	4	US-10-184-638-236	Sequence 236, App
395	31	66.0	284	4	US-10-197-658-236	Sequence 236, App	468	31	66.0	284	4	US-10-187-752-236	Sequence 236, App
396	31	66.0	284	4	US-10-195-894-236	Sequence 236, App	469	31	66.0	284	4	US-10-187-887-236	Sequence 236, App
397	31	66.0	284	4	US-10-176-484-236	Sequence 236, App	470	31	66.0	284	4	US-10-194-461-236	Sequence 236, App
398	31	66.0	284	4	US-10-176-753-236	Sequence 236, App	471	31	66.0	284	4	US-10-195-892-236	Sequence 236, App
399	31	66.0	284	4	US-10-176-917-236	Sequence 236, App	472	31	66.0	284	4	US-10-196-751-236	Sequence 236, App
400	31	66.0	284	4	US-10-176-982-236	Sequence 236, App	473	31	66.0	284	4	US-10-197-694-236	Sequence 236, App
401	31	66.0	284	4	US-10-179-506-236	Sequence 236, App	474	31	66.0	284	4	US-10-197-697-236	Sequence 236, App
402	31	66.0	284	4	US-10-179-513-236	Sequence 236, App	475	31	66.0	284	4	US-10-197-707-236	Sequence 236, App
403	31	66.0	284	4	US-10-179-514-236	Sequence 236, App	476	31	66.0	284	4	US-10-199-318-236	Sequence 236, App
404	31	66.0	284	4	US-10-179-522-236	Sequence 236, App	477	31	66.0	284	4	US-10-199-411-236	Sequence 236, App
405	31	66.0	284	4	US-10-180-556-236	Sequence 236, App	478	31	66.0	284	4	US-10-199-458-236	Sequence 236, App
406	31	66.0	284	4	US-10-180-560-236	Sequence 236, App	479	31	66.0	284	4	US-10-199-462-236	Sequence 236, App
407	31	66.0	284	4	US-10-183-015-236	Sequence 236, App	480	31	66.0	284	4	US-10-201-324-236	Sequence 236, App
408	31	66.0	284	4	US-10-184-615-236	Sequence 236, App	481	31	66.0	284	4	US-10-201-328-236	Sequence 236, App
409	31	66.0	284	4	US-10-184-620-236	Sequence 236, App	482	31	66.0	284	4	US-10-201-527-236	Sequence 236, App
410	31	66.0	284	4	US-10-184-643-236	Sequence 236, App	483	31	66.0	284	4	US-10-201-528-236	Sequence 236, App
411	31	66.0	284	4	US-10-184-656-236	Sequence 236, App	484	31	66.0	284	4	US-10-201-529-236	Sequence 236, App
412	31	66.0	284	4	US-10-192-010-236	Sequence 236, App	485	31	66.0	284	4	US-10-201-530-236	Sequence 236, App
413	31	66.0	284	4	US-10-063-553-62	Sequence 62, Appl	486	31	66.0	284	4	US-10-202-408-236	Sequence 236, App
414	31	66.0	284	4	US-10-205-908-236	Sequence 236, App	487	31	66.0	284	4	US-10-202-409-236	Sequence 236, App
415	31	66.0	284	4	US-10-063-518-62	Sequence 62, Appl	488	31	66.0	284	4	US-10-202-411-236	Sequence 236, App
416	31	66.0	284	4	US-10-184-619-236	Sequence 236, App	489	31	66.0	284	4	US-10-202-472-236	Sequence 236, App
417	31	66.0	284	4	US-10-187-599-236	Sequence 236, App	490	31	66.0	284	4	US-10-205-502-236	Sequence 236, App
418	31	66.0	284	4	US-10-187-750-236	Sequence 236, App	491	31	66.0	284	4	US-10-205-507-236	Sequence 236, App
419	31	66.0	284	4	US-10-188-780-236	Sequence 236, App	492	31	66.0	284	4	US-10-205-511-236	Sequence 236, App
420	31	66.0	284	4	US-10-192-015-236	Sequence 236, App	493	31	66.0	284	4	US-10-205-902-236	Sequence 236, App
421	31	66.0	284	4	US-10-194-394-236	Sequence 236, App	494	31	66.0	284	4	US-10-205-907-236	Sequence 236, App
422	31	66.0	284	4	US-10-194-425-236	Sequence 236, App	495	31	66.0	284	4	US-10-176-484-236	Sequence 236, App
423	31	66.0	284	4	US-10-194-485-236	Sequence 236, App	496	31	66.0	284	4	US-10-194-456-236	Sequence 236, App
424	31	66.0	284	4	US-10-195-885-236	Sequence 236, App	497	31	66.0	284	4	US-10-196-758-236	Sequence 236, App
425	31	66.0	284	4	US-10-195-899-236	Sequence 236, App	498	31	66.0	284	4	US-10-198-770-236	Sequence 236, App
426	31	66.0	284	4	US-10-196-748-236	Sequence 236, App	499	31	66.0	284	4	US-10-199-308-236	Sequence 236, App
427	31	66.0	284	4	US-10-196-750-236	Sequence 236, App	500	31	66.0	284	4	US-10-200-617-236	Sequence 236, App
428	31	66.0	284	4	US-10-197-699-236	Sequence 236, App	501	31	66.0	284	4	US-10-205-893-236	Sequence 236, App
429	31	66.0	284	4	US-10-197-700-236	Sequence 236, App	502	31	66.0	284	4	US-10-205-897-236	Sequence 236, App
430	31	66.0	284	4	US-10-197-705-236	Sequence 236, App	503	31	66.0	284	4	US-10-063-563-62	Sequence 62, Appl
431	31	66.0	284	4	US-10-197-708-236	Sequence 236, App	504	31	66.0	284	4	US-10-195-896-236	Sequence 236, App
432	31	66.0	284	4	US-10-198-764-236	Sequence 236, App	505	31	66.0	284	4	US-10-180-550-236	Sequence 236, App
433	31	66.0	284	4	US-10-198-765-236	Sequence 236, App	506	31	66.0	284	4	US-10-183-014-236	Sequence 236, App
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435	31	66.0	284	4	US-10-198-769-236	Sequence 236, App	508	31	66.0	284	4	US-10-187-740-236	Sequence 236, App
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ALIGNMENTS

RESULT 1
US-10-476-570-20 Application US/10476570
; Sequence 20. Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 20
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 17-31
US-10-476-570-20

Query Match 100.0%; Score 47; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
Db 5 QLCLEQTT 13

RESULT 2
US-10-476-570-21 Application US/10476570
; Sequence 21. Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:

APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 21
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 20-34
US-10-476-570-21

Query Match 100.0%; Score 47; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
Db 2 QLCLEQTT 10

RESULT 3
US-10-476-570-8 Application US/10476570
; Sequence 8. Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 8
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-34
US-10-476-570-8

Query Match 100.0%; Score 47; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QLCLEQTT 9
Db 8 QLCLEQTT 16


```
RESULT 4
US-11-021-949-1
; Sequence 1, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SABIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-1
```

```
Query Match          100.0%; Score 47; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.17;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 QUCTELQTT 9
        |||||
        13 QUCTELQTT 21
```

```
RESULT 5
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53
```

```
Query Match          100.0%; Score 47; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 QUCTELQTT 9
        |||||
        7 QUCTELQTT 15
```

```
RESULT 6
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4
```

```
Query Match          100.0%; Score 47; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 QUCTELQTT 9
        |||||
        7 QUCTELQTT 15
```

```
RESULT 7
US-10-476-570-9
; Sequence 9, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 32
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9
```

```
Query Match          100.0%; Score 47; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 QUCTELQTT 9
```

Db 8 QCTELQTT 16
|||||

RESULT 8
US-10-476-570-19

; Sequence 19, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUEVILLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 33
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19

Query Match 100.0%; Score 47; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.22;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
|||||
Db 8 QCTELQTT 16

RESULT 9

US-10-177-390-6
; Sequence 6, Application US/10177390
; Publication No. US20030143743A1
; GENERAL INFORMATION:
; APPLICANT: Schuler, Gerold
; APPLICANT: N.V. Antwerp Innovatiecentrum
; TITLE OF INVENTION: Improved Transfection of Eucaryotic Cells with linear
; FILE REFERENCE: 021505wo/JH/ml
; CURRENT APPLICATION NUMBER: US/10/177,390
; CURRENT FILING DATE: 2002-06-20
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match 100.0%; Score 47; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.99;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
|||||
Db 14 QCTELQTT 22

. RESULT 10

US-10-484-063-20
; Sequence 20, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILTERMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-20

Query Match 100.0%; Score 47; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.99;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
|||||
Db 14 QCTELQTT 22

RESULT 11

US-10-484-063-27
; Sequence 27, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILTERMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 27
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match 100.0%; Score 47; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.99;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
|||||
Db 14 QCTELQTT 22

RESULT 12

US-10-858-384-2
; Sequence 2, Application US/10858384
; Publication No. US20050033025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE

```

; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCES
; APPLICANT: PERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match          100.0%; Score 47; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELOT 9
DB 21 QCTELOT 29

RESULT 13
US-10-367-057-16
; Sequence 16, Application US/10367057
; Publication No. US20050100554A1
; GENERAL INFORMATION:
; APPLICANT: Cuthill, Scott;
; APPLICANT: JACKSON, Amanda;
; APPLICANT: Lewin, David A.;
; APPLICANT: Ooi, Chean Eng
; TITLE OF INVENTION: Complexes and Methods of Using Same
; FILE REFERENCE: 21402-559
; CURRENT APPLICATION NUMBER: US/10/367,057
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: 60/256,911
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 198
; SOFTWARE: CuraSeqList version 0.1
; SEQ ID NO 16
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-367-057-16

Query Match          100.0%; Score 47; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELOT 9
DB 21 QCTELOT 29

RESULT 14
US-11-021-949-13
; Sequence 13, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
```

```

; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match          100.0%; Score 47; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELOT 9
DB 21 QCTELOT 29

RESULT 15
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; CURRENT FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03271
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match          100.0%; Score 47; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELOT 9
DB 26 QCTELOT 34

RESULT 16
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOU, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; CURRENT FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
```

; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone B6*TMF.
US-11-072-288-1

Query Match 100.0%; Score 47; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 49 QLCLEQTT 57

RESULT 17
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Publication No. US20020081329A1
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; PRIOR FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 47; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 1.7;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 21 QLCLEQTT 29

RESULT 18
US-10-000-903-4
; Sequence 4, Application US/10000903
; Publication No. US2002018221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; PRIOR FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22

; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-4

Query Match 100.0%; Score 47; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 127 QLCLEQTT 135

RESULT 19
US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; PRIOR FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4

Query Match 100.0%; Score 47; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.8;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 127 QLCLEQTT 135

RESULT 20
US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US2002018221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; PRIOR FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5

PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO: 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-10

Query Match 100.0%; Score 47; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.9; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELQTT 9
Db 146 QCTELQTT 154

RESULT 21
US-10-899-771-10
Sequence 10, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO: 10
LENGTH: 292
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimaeric protein (Clyra from Streptococcus
OTHER INFORMATION: pneumoniae and B6 from Human papilloma virus type
OTHER INFORMATION: 16)
US-10-899-771-10

Query Match 100.0%; Score 47; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.9; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELQTT 9
Db 146 QCTELQTT 154

RESULT 22
US-10-000-903-6
Sequence 6, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17

PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO: 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-6

Query Match 100.0%; Score 47; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 2.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELQTT 9
Db 127 QCTELQTT 135

RESULT 23
US-10-899-771-6
Sequence 6, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO: 6
LENGTH: 371
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimaeric protein (protein D from Haemophilus
OTHER INFORMATION: influenzae B and B6B7 fusion from Human papilloma
OTHER INFORMATION: virus type 16)
US-10-899-771-6

Query Match 100.0%; Score 47; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 2.4; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELQTT 9
Db 127 QCTELQTT 135

RESULT 24
US-10-000-903-14
Sequence 14, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285

;; PRIOR FILING DATE: 1998-08-17
;; PRIOR APPLICATION NUMBER: GB 9717953.5
;; PRIOR FILING DATE: 1997-08-22
;; NUMBER OF SEQ ID NOS: 23
;; SOFTWARE: FastSeq for Windows Version 3.0
;; SEQ ID NO 14
;; LENGTH: 390
;; TYPE: PRT
;; ORGANISM: Homo sapien
US-10-000-903-14

Query Match 100.0%; Score 47; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 146 QLCLEQTT 154

RESULT 25
US-10-899-771-14
;; Sequence 14, Application US/10899771
;; Publication No. US20050031638A1
;; GENERAL INFORMATION:
;; APPLICANT: Dalemans, Wilfried L.J.
;; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
;; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
;; FILE REFERENCE: B45124
;; CURRENT APPLICATION NUMBER: US/10/899,771
;; CURRENT FILING DATE: 2004-07-27
;; PRIOR APPLICATION NUMBER: US/09/581,976
;; PRIOR FILING DATE: 2000-06-20
;; PRIOR APPLICATION NUMBER: PCT/EP98/08563
;; PRIOR FILING DATE: 1998-12-18
;; PRIOR APPLICATION NUMBER: GB 9727262.9
;; PRIOR FILING DATE: 1997-12-24
;; NUMBER OF SEQ ID NOS: 28
;; SOFTWARE: FastSeq for Windows Version 3.0
;; SEQ ID NO 14
;; LENGTH: 390
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: Chimeraic protein (Clyta from Streptococcus
;; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
;; OTHER INFORMATION: virus type 16)
US-10-899-771-14

Query Match 100.0%; Score 47; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 146 QLCLEQTT 154

RESULT 26
US-10-367-095-10
;; Sequence 10, Application US/10367095
;; Publication No. US20030228696A1
;; GENERAL INFORMATION:
;; APPLICANT: Robin A. Robinson
;; TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
;; FILE REFERENCE: 44149-1US1
;; CURRENT APPLICATION NUMBER: US/10/367,095
;; CURRENT FILING DATE: 2003-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14

;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,135
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-095-10

Query Match 100.0%; Score 47; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QLCLEQTT 9
Db 491 QLCLEQTT 499

RESULT 27
US-10-368-046-10
;; Sequence 10, Application US/10368046
;; Publication No. US20040063188A1
;; GENERAL INFORMATION:
;; APPLICANT: Robin A. Robinson
;; TITLE OF INVENTION: Method for Isolation and Purification of
;; TITLE OF INVENTION: Expressed Gene Products In Vitro
;; FILE REFERENCE: 44149-3US1
;; CURRENT APPLICATION NUMBER: US/10/368,046
;; CURRENT FILING DATE: 2003-02-15
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,135
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536

```

; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-368-046-10

Query Match          100.0%; Score 47; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 491 QCTELQTT 499

RESULT 28
US-10-367-367-10
; Sequence 10, Application US/10367367
; Publication No. US20040121465A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 44149-2US1
; CURRENT APPLICATION NUMBER: US/10/367,367
; PRIOR FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-367-10

Query Match          100.0%; Score 47; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 491 QCTELQTT 499

RESULT 29
US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:
; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 19065/2132
; CURRENT APPLICATION NUMBER: US/10/918,337
; PRIOR FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
```

```

; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10

Query Match          100.0%; Score 47; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 491 QCTELQTT 499

RESULT 30
US-09-925-299-1386
; Sequence 1386, Application US/09925299
; Patent No. US20020055627A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA102
; CURRENT APPLICATION NUMBER: US/09/925,299
; PRIOR FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05883
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 1556
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1386
; LENGTH: 48
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (5)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (13)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (15)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (40)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-299-1386

Query Match          78.7%; Score 37; DB 3; Length 48;
Best Local Similarity 87.5%; Pred. No. 20;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
```

OY 2 LCTELQTT 9
Db 26 LCTELQTT 33

RESULT 31

US-09-925-299-1386
; Sequence 1386, Application US/09925299
; Publication No. US20030040617A9
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA102
; CURRENT APPLICATION NUMBER: US/09/925,299
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05883
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; NUMBER OF SEQ ID NOS: 1556
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1386
; LENGTH: 48
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (5)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (13)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (15)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (40)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-299-1386

Query Match 78.7%; Score 37; DB 3; Length 48;
Best Local Similarity 87.5%; Pred. No. 20;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

OY 2 LCTELQTT 9
Db 26 LCTELQTT 33

RESULT 32

US-10-437-963-115784
; Sequence 115784, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 115784
; LENGTH: 587
; TYPE: PRT
; ORGANISM: Oryza sativa

FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(587)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_19348C.1.pcp
US-10-437-963-115784

Query Match 78.7%; Score 37; DB 4; Length 587;
Best Local Similarity 77.8%; Pred. No. 2,4e+02;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 1 QLCGELQTT 9
Db 457 QLCGELQTT 465

RESULT 33

US-10-437-963-122232
; Sequence 122232, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 122232
; LENGTH: 853
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(853)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_2517C.1.pcp
US-10-437-963-122232

Query Match 78.7%; Score 37; DB 4; Length 853;
Best Local Similarity 66.7%; Pred. No. 3,5e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 QLCGELQTT 9
Db 753 QLCGELQTT 761

RESULT 34

US-10-424-599-191823
; Sequence 191823, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 191823


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/ LENGTH: 55
/ TYPE: PRT
/ ORGANISM: Glycine max
/ FEATURE:
/ OTHER INFORMATION: Clone ID: PAT_MRT3847_15236C.1.pcp
US-10-424-599-191823

Query Match          76.6%; Score 36; DB 4; Length 55;
Best Local Similarity 87.5%; Pred. No. 35;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 13 LCMELQTT 20

RESULT 35
US-10-369-493-1712
/ Sequence 1712, Application US/10369493
/ Publication No. US20030233675A1
/ GENERAL INFORMATION:
/ APPLICANT: Cao, Yongwei
/ APPLICANT: Hinkle, Gregory J.
/ APPLICANT: Slater, Steven C.
/ APPLICANT: Goldman, Barry S.
/ APPLICANT: Chen, Xianfeng
/ TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
/ FILE REFERENCE: 38-10(52052)B
/ CURRENT APPLICATION NUMBER: US/10/369,493
/ PRIOR FILING DATE: 2003-02-28
/ PRIOR APPLICATION NUMBER: US 60/360,039
/ PRIOR FILING DATE: 2002-02-21
/ NUMBER OF SEQ ID NOS: 47374
/ SEQ ID NO 1712
/ LENGTH: 470
/ TYPE: PRT
/ ORGANISM: Saccharomyces cerevisiae
US-10-369-493-1712

Query Match          76.6%; Score 36; DB 4; Length 470;
Best Local Similarity 87.5%; Pred. No. 3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 1 QCTELQTT 8
Db 357 QLCBELQTT 364

RESULT 36
US-10-282-122A-70249
/ Sequence 70249, Application US/10282122A
/ Publication No. US20040029129A1
/ GENERAL INFORMATION:
/ APPLICANT: Wang, Liangsu
/ APPLICANT: Zamudio, Carlos
/ APPLICANT: Malone, Cheryl
/ APPLICANT: Haeselbeck, Robert
/ APPLICANT: Ohlsen, Karl
/ APPLICANT: Zykkind, Judith
/ APPLICANT: Wall, Daniel
/ APPLICANT: Trawick, John
/ APPLICANT: Carr, Grant
/ APPLICANT: Yamamoto, Robert
/ APPLICANT: Forsyth, R.
/ APPLICANT: Xu, H.
/ TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
/ FILE REFERENCE: ELITRA.034A
/ CURRENT FILING DATE: 2003-02-20
/ PRIOR APPLICATION NUMBER: US/10/282,122A
/ PRIOR FILING DATE: 2000-03-21
/ PRIOR APPLICATION NUMBER: 60/206,848
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/ PRIOR FILING DATE: 2000-05-23
/ PRIOR APPLICATION NUMBER: 60/207,727
/ PRIOR FILING DATE: 2000-05-26
/ PRIOR APPLICATION NUMBER: 60/230,335
/ PRIOR FILING DATE: 2000-09-06
/ PRIOR APPLICATION NUMBER: 60/230,347
/ PRIOR FILING DATE: 2000-09-09
/ PRIOR APPLICATION NUMBER: 60/242,578
/ PRIOR FILING DATE: 2000-10-23
/ PRIOR APPLICATION NUMBER: 60/253,625
/ PRIOR FILING DATE: 2000-11-27
/ PRIOR APPLICATION NUMBER: 60/257,931
/ PRIOR FILING DATE: 2000-12-22
/ PRIOR APPLICATION NUMBER: 60/267,636
/ PRIOR FILING DATE: 2001-02-09
/ PRIOR APPLICATION NUMBER: 60/269,308
/ PRIOR FILING DATE: 2001-02-16
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 78614
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 70249
/ LENGTH: 158
/ TYPE: PRT
/ ORGANISM: Staphylococcus aureus
US-10-282-122A-70249

Query Match          74.5%; Score 35; DB 4; Length 158;
Best Local Similarity 75.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 57 LCMELQTT 64

RESULT 37
US-10-857-625-567
/ Sequence 567, Application US/10857625
/ Publication No. US20050026189A1
/ GENERAL INFORMATION:
/ APPLICANT: Wang, Liangsu
/ APPLICANT: Zamudio, Carlos
/ TITLE OF INVENTION: MICROBIAL OPERONS
/ FILE REFERENCE: ELITRA.036A
/ CURRENT APPLICATION NUMBER: US/10/857,625
/ PRIOR FILING DATE: 2004-05-28
/ PRIOR APPLICATION NUMBER: 60/474768
/ PRIOR FILING DATE: 2003-05-29
/ NUMBER OF SEQ ID NOS: 833
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 567
/ LENGTH: 158
/ TYPE: PRT
/ ORGANISM: Staphylococcus aureus
US-10-857-625-567

Query Match          74.5%; Score 35; DB 5; Length 158;
Best Local Similarity 75.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 57 LCMELQTT 64

RESULT 38
US-10-820-155-14
/ Sequence 14, Application US/10820155
/ Publication No. US20050137126A1
/ GENERAL INFORMATION:
/ APPLICANT: Natimmune A/S
/ APPLICANT: Wellguny, Dietmar
/ APPLICANT: Jensenius, Jens Christian
```

```

; APPLICANT: Kongerslev, Lelf
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 14
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-14

```

```

Query Match      74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

Qy      2 LCTELOTT 9
        |||||
Db      145 LCTELOGT 152

```

```

RESULT 39
US-10-820-155-23
; Sequence 23, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Weilguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Lelf
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 23
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-23

```

```

Query Match      74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

Qy      2 LCTELOTT 9
        |||||
Db      145 LCTELOGT 152

```

```

RESULT 40
US-10-820-155-30
; Sequence 30, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Weilguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Lelf
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 30
; LENGTH: 239
; TYPE: PRT

```

```

; ORGANISM: Mus musculus
US-10-820-155-30

```

```

Query Match      74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

Qy      2 LCTELOTT 9
        |||||
Db      145 LCTELOGT 152

```

```

RESULT 41
US-10-820-155-39
; Sequence 39, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Weilguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Lelf
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 39
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-39

```

```

Query Match      74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

Qy      2 LCTELOTT 9
        |||||
Db      145 LCTELOGT 152

```

```

RESULT 42
US-10-820-155-85
; Sequence 85, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Weilguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Lelf
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 85
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-85

```

```

Query Match      74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

Qy      2 LCTELOTT 9
        |||||
Db      145 LCTELOGT 152

```

```
RESULT 43
US-10-820-155-94
; Sequence 94, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Wellguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; CURRENT FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 94
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: MISC_FEATURE
; LOCATION: (91)..(116)
; OTHER INFORMATION: Unknown
US-10-820-155-94

Query Match          74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 145 LCTELQTT 152

RESULT 44
US-10-820-155-97
; Sequence 97, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: NatImmune A/S
; APPLICANT: Wellguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; CURRENT FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 97
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-97

Query Match          74.5%; Score 35; DB 5; Length 239;
Best Local Similarity 87.5%; Pred. No. 2.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 145 LCTELQTT 152

RESULT 45
US-10-424-599-246277
; Sequence 246277, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J

; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 246277
; LENGTH: 104
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_6441C.1.pep
US-10-424-599-246277

Query Match          72.3%; Score 34; DB 4; Length 104;
Best Local Similarity 66.7%; Pred. No. 1.5e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 QLCTELQTT 9
Db 8 ELCTKLQTT 16

RESULT 46
US-10-767-701-37218
; Sequence 37218, Application US/10767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53535)B
; CURRENT APPLICATION NUMBER: US/10/767,701
; CURRENT FILING DATE: 2004-01-29
; NUMBER OF SEQ ID NOS: 63128
; SEQ ID NO 37218
; LENGTH: 120
; TYPE: PRT
; ORGANISM: Sorghum bicolor
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(120)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: SORBI-28MAY03-C2069_1.pep
US-10-767-701-37218

Query Match          72.3%; Score 34; DB 4; Length 120;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 111 LCTELQTT 118

RESULT 47
US-10-127-816-9
; Sequence 9, Application US/10127816
; Publication No. US20030104416A1
; GENERAL INFORMATION:
; APPLICANT: Sheperd, Paul O.
; APPLICANT: Fox, Brian A.
; APPLICANT: Klucher, Kevin M.
; APPLICANT: Taft, David W.
; APPLICANT: Kindvogel, Wayne R.
; TITLE OF INVENTION: CYTOKINE PROTEIN FAMILY
; FILE REFERENCE: 01-17
```

```

; CURRENT APPLICATION NUMBER: US/10/127,816
; CURRENT FILING DATE: 2002-04-19
; PRIOR APPLICATION NUMBER: US 60/285,408
; PRIOR FILING DATE: 2001-04-20
; PRIOR APPLICATION NUMBER: US 60/286,482
; PRIOR FILING DATE: 2001-04-25
; PRIOR APPLICATION NUMBER: US 60/341,050
; PRIOR FILING DATE: 2001-10-22
; PRIOR APPLICATION NUMBER: US 60/341,105
; PRIOR FILING DATE: 2001-10-22
; PRIOR APPLICATION NUMBER: US 09/895,834
; PRIOR FILING DATE: 2001-06-29
; PRIOR APPLICATION NUMBER: US 60/285,424
; PRIOR FILING DATE: 2001-04-20
; NUMBER OF SEQ ID NOS: 59
; SOFTWARE: FaestSeq for Windows Version 4.0
; SEQ ID NO 9
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-127-816-9
```

```

Query Match          72.3%; Score 34; DB 4; Length 202;
Best Local Similarity 66.7%; Pred. No. 3e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 QCTCTLOQT 9
      |||:|
Db      139 QCTCTLOQT 147
```

```

RESULT 48
US-10-127-816-11
; Sequence 11, Application US/10127816
; Publication No. US20030104416A1
; GENERAL INFORMATION:
; APPLICANT: Sheppard, Paul O.
; APPLICANT: Fox, Brian A.
; APPLICANT: Klucher, Kevin M.
; APPLICANT: Taft, David W.
; APPLICANT: Kindsvogel, Wayne R.
; TITLE OF INVENTION: CYTOKINE PROTEIN FAMILY
; FILE REFERENCE: 01-17
; CURRENT APPLICATION NUMBER: US/10/127,816
; CURRENT FILING DATE: 2002-04-19
; PRIOR APPLICATION NUMBER: US 60/285,408
; PRIOR FILING DATE: 2001-04-20
; PRIOR APPLICATION NUMBER: US 60/286,482
; PRIOR FILING DATE: 2001-04-25
; PRIOR APPLICATION NUMBER: US 60/341,050
; PRIOR FILING DATE: 2001-10-22
; PRIOR APPLICATION NUMBER: US 60/341,105
; PRIOR FILING DATE: 2001-10-22
; PRIOR APPLICATION NUMBER: US 09/895,834
; PRIOR FILING DATE: 2001-06-29
; PRIOR APPLICATION NUMBER: US 60/285,424
; PRIOR FILING DATE: 2001-04-20
; NUMBER OF SEQ ID NOS: 59
; SOFTWARE: FaestSeq for Windows Version 4.0
; SEQ ID NO 11
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-127-816-11
```

```

Query Match          72.3%; Score 34; DB 4; Length 202;
Best Local Similarity 66.7%; Pred. No. 3e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 QCTCTLOQT 9
      |||:|
Db      139 QCTCTLOQT 147
```

```

RESULT 49
US-10-142-717-12
; Sequence 12, Application US/10142717
; Publication No. US20030104579A1
; GENERAL INFORMATION:
; APPLICANT: Immunex Corporation
; APPLICANT: Baum, Peter R
; APPLICANT: Mosley, Bruce A
; APPLICANT: Ketchem, Randal R
; APPLICANT: Taylor, Scott L
; TITLE OF INVENTION: CYTOKINE POLYPEPTIDES
; FILE REFERENCE: 3282-A
; CURRENT APPLICATION NUMBER: US/10/142,717
; CURRENT FILING DATE: 2002-05-08
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 12
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-142-717-12
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```

Query Match          72.3%; Score 34; DB 4; Length 202;
Best Local Similarity 66.7%; Pred. No. 3e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 QCTCTLOQT 9
      |||:|
Db      139 QCTCTLOQT 147
```

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RESULT 50
US-10-420-034A-60
; Sequence 60, Application US/10420034A
; Publication No. US20040029228A1
; GENERAL INFORMATION:
; APPLICANT: Presnell, Scott R.
; APPLICANT: Xu, Wenfeng
; APPLICANT: No. US20040029228A1ak, Julia E.
; APPLICANT: Whitmore, Theodore E.
; APPLICANT: Grant, Francis J.
; APPLICANT: Kindsvogel, Wayne R.
; APPLICANT: Klucher, Kevin M.
; TITLE OF INVENTION: CYTOKINE RECEPTOR
; FILE REFERENCE: 02-10
; CURRENT APPLICATION NUMBER: US/10/420,034A
; CURRENT FILING DATE: 2003-04-18
; PRIOR APPLICATION NUMBER: 60/373,813
; PRIOR FILING DATE: 2002-04-19
; NUMBER OF SEQ ID NOS: 69
; SOFTWARE: FaestSeq for Windows Version 4.0
; SEQ ID NO 60
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-420-034A-60
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```

Query Match          72.3%; Score 34; DB 4; Length 202;
Best Local Similarity 66.7%; Pred. No. 3e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
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QY      1 QCTCTLOQT 9
      |||:|
Db      139 QCTCTLOQT 147
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Search completed: May 5, 2006, 08:50:44
Job time : 59.3 secs
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GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:40:52 ; Search time 8.4 Seconds
(without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-3
Perfect score: 47
Sequence: 1 QLTCLQTT 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

Published Applications AA_New:*
1: /SID5/pcodata/1/pubppa/US08_NEW_PUB.pep1:*
2: /SID5/pcodata/1/pubppa/US06_NEW_PUB.pep:*
3: /SID5/pcodata/1/pubppa/US07_NEW_PUB.pep:*
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5: /SID5/pcodata/1/pubppa/PCT_NEW_PUB.pep:*
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12: /SID5/pcodata/1/pubppa/US60_NEW_PUB.pep1:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	47	100.0	151	9	US-10-530-253-13
2	47	100.0	158	11	US-11-206-138-3
3	47	100.0	248	9	US-10-530-253-1
4	47	100.0	248	9	US-10-530-253-3
5	47	100.0	248	9	US-10-530-253-5
6	47	100.0	248	9	US-10-530-253-7
7	47	100.0	248	9	US-10-530-253-9
8	47	100.0	248	9	US-10-530-253-11
9	47	100.0	256	11	US-11-192-923A-2
10	47	100.0	252	11	US-11-098-662-8
11	47	100.0	252	11	US-11-098-662-10
12	47	100.0	252	11	US-11-193-955-8
13	47	100.0	252	11	US-11-193-955-10
14	47	100.0	252	11	US-11-266-626-9
15	47	100.0	252	11	US-11-266-626-11
16	47	100.0	252	11	US-10-530-253-15
17	47	100.0	252	11	US-10-530-253-20
18	47	100.0	252	11	US-11-079-463-6325
19	47	100.0	252	11	US-11-087-099-6055
20	47	100.0	252	11	US-11-072-512-2475
21	47	100.0	252	11	US-10-467-657-7248

22	31	66.0	129	9	US-10-519-390-15	Sequence 15, App1
23	31	66.0	129	11	US-11-176-830-207	Sequence 207, App
24	31	66.0	129	11	US-11-176-830-546	Sequence 546, App
25	31	66.0	129	11	US-11-176-830-547	Sequence 547, App
26	31	66.0	129	11	US-11-176-830-548	Sequence 548, App
27	31	66.0	129	11	US-11-176-830-549	Sequence 549, App
28	31	66.0	129	11	US-11-176-830-550	Sequence 550, App
29	31	66.0	129	11	US-11-176-830-551	Sequence 551, App
30	31	66.0	129	11	US-11-176-830-552	Sequence 552, App
31	31	66.0	129	11	US-11-176-830-553	Sequence 553, App
32	31	66.0	129	11	US-11-176-830-554	Sequence 554, App
33	31	66.0	129	11	US-11-176-830-555	Sequence 555, App
34	31	66.0	129	11	US-11-176-830-556	Sequence 556, App
35	31	66.0	129	11	US-11-176-830-557	Sequence 557, App
36	31	66.0	129	11	US-11-176-830-558	Sequence 558, App
37	31	66.0	129	11	US-11-176-830-559	Sequence 559, App
38	31	66.0	129	11	US-11-176-830-560	Sequence 560, App
39	31	66.0	129	11	US-11-176-830-561	Sequence 561, App
40	31	66.0	129	11	US-11-176-830-562	Sequence 562, App
41	31	66.0	129	11	US-11-176-830-563	Sequence 563, App
42	31	66.0	129	11	US-11-176-830-564	Sequence 564, App
43	31	66.0	129	11	US-11-176-830-565	Sequence 565, App
44	31	66.0	129	11	US-11-176-830-566	Sequence 566, App
45	31	66.0	129	11	US-11-176-830-567	Sequence 567, App
46	31	66.0	150	9	US-10-469-561-25	Sequence 25, App1
47	31	66.0	153	8	US-10-511-937-2476	Sequence 2476, App
48	31	66.0	153	11	US-11-174-398-8	Sequence 8, App1
49	31	66.0	153	11	US-11-289-226-13	Sequence 19, App1
50	31	66.0	155	9	US-10-530-253-19	Sequence 5, App1
51	31	66.0	258	11	US-11-026-396-5	Sequence 7, App1
52	31	66.0	273	9	US-10-063-703-62	Sequence 62, App1
53	31	66.0	284	9	US-10-194-447-236	Sequence 236, App
54	31	66.0	284	9	US-10-195-883-236	Sequence 236, App
55	31	66.0	284	9	US-10-195-888-236	Sequence 236, App
56	31	66.0	284	9	US-10-195-889-236	Sequence 236, App
57	31	66.0	284	11	US-11-102-240-62	Sequence 62, App1
58	31	66.0	284	11	US-11-103-195-62	Sequence 62, App1
59	31	66.0	293	11	US-11-026-396-4	Sequence 4, App1
60	31	66.0	293	11	US-11-026-396-2	Sequence 2, App1
61	31	66.0	295	11	US-11-087-099-5930	Sequence 5930, App
62	31	66.0	400	11	US-11-188-298-22075	Sequence 22075, A
63	31	66.0	412	11	US-11-087-099-2684	Sequence 2684, App
64	31	66.0	492	11	US-11-096-568A-14735	Sequence 14735, A
65	31	66.0	492	11	US-11-131-479-2	Sequence 2, App1
66	31	66.0	498	11	US-11-131-479-76	Sequence 76, App1
67	31	66.0	500	11	US-11-096-568A-14734	Sequence 14734, A
68	31	66.0	514	11	US-11-087-099-2712	Sequence 2712, App1
69	31	66.0	522	11	US-11-131-479-9	Sequence 9, App1
70	31	66.0	522	11	US-11-131-479-7	Sequence 7, App1
71	31	66.0	522	11	US-11-131-479-9	Sequence 9, App1
72	31	66.0	587	11	US-11-168-288-296	Sequence 296, App
73	31	66.0	587	11	US-11-168-288-296	Sequence 296, App
74	31	66.0	146	11	US-11-096-568A-14842	Sequence 14842, A
75	30	63.8	172	11	US-11-177-010-4	Sequence 4, App1
76	30	63.8	172	11	US-11-112-369-1	Sequence 1, App1
77	30	63.8	172	11	US-11-112-369-2	Sequence 2, App1
78	30	63.8	172	11	US-11-112-369-2	Sequence 2, App1
79	30	63.8	172	11	US-11-298-955-2	Sequence 3, App1
80	30	63.8	172	11	US-11-298-955-3	Sequence 3, App1
81	30	63.8	174	11	US-11-087-099-1876	Sequence 1876, App
82	30	63.8	175	11	US-11-087-099-9020	Sequence 9020, App
83	30	63.8	175	11	US-11-087-099-9020	Sequence 9020, App
84	30	63.8	231	11	US-11-096-568A-3393	Sequence 3393, App
85	30	63.8	277	8	US-10-511-937-2455	Sequence 2455, App
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93	30	63.8	376	11	US-11-188-298-11484	Sequence 11484, App
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102	29	61.7	492	11	US-11-264-096-1232	Sequence 1232, Ap
103	29	61.7	499	11	US-11-096-568A-6201	Sequence 6201, Ap
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107	29	61.7	834	9	US-10-501-035-280	Sequence 280, App
108	29	61.7	851	9	US-10-330-773-397	Sequence 397, App
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155	28	59.6	128	9	US-10-475-075-481	Sequence 481, App
156	28	59.6	129	11	US-11-045-004-977	Sequence 977, App
157	28	59.6	129	11	US-11-045-004-977	Sequence 977, App
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159	28	59.6	146	11	US-11-096-568A-29968	Sequence 29968, A
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162	28	59.6	148	11	US-11-075-047A-89	Sequence 89, Appl
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168	28	59.6	231	9	US-10-915-002-351	Sequence 351, App
169	28	59.6	269	11	US-11-096-568A-5631	Sequence 5631, Ap
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179	28	59.6	368	11	US-11-087-099-4532	Sequence 4532, Ap
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182	28	59.6	413	8	US-10-511-937-2428	Sequence 2428, Ap
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185	28	59.6	446	11	US-11-079-463-5912	Sequence 5912, Ap
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187	28	59.6	480	11	US-11-096-568A-19197	Sequence 19197, A
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191	28	59.6	589	9	US-10-880-881-49	Sequence 49, Appl
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195	28	59.6	752	9	US-10-506-454-648	Sequence 648, App
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208	28	59.6	941	11	US-11-124-367A-315	Sequence 315, App
209	28	59.6	1240	9	US-10-501-035-260	Sequence 260, App
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219	28	59.6	1883	11	US-11-126-022-14	Sequence 14, Appl
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222	28	59.6	1883	11	US-11-126-022-17	Sequence 17, Appl
223	28	59.6	1883	11	US-11-126-022-18	Sequence 18, Appl
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225	28	59.6	1883	11	US-11-126-022-20	Sequence 20, Appl
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237	27	57.4	10	11	US-11-045-024-9684	Sequence 9684, Ap
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244	27	57.4	11	11	US-11-045-024-11853	Sequence 11853, A	317	27	57.4	561	11	US-11-029-465-12	Sequence 10, Appl1
245	27	57.4	15	11	US-11-045-024-13252	Sequence 13252, A	318	27	57.4	562	9	US-10-507-928-10	Sequence 10, Appl1
246	27	57.4	15	11	US-11-045-024-13433	Sequence 13433, A	319	27	57.4	562	9	US-10-507-928-12	Sequence 12, Appl1
247	27	57.4	19	11	US-11-004-399-1357	Sequence 1357, Ap	320	27	57.4	562	11	US-11-029-465-10	Sequence 10, Appl1
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249	27	57.4	34	11	US-11-207-078-256	Sequence 256, App	322	27	57.4	565	11	US-11-096-568A-1993	Sequence 1993, A
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252	27	57.4	69	11	US-11-087-099-4917	Sequence 4917, Ap	325	27	57.4	625	11	US-11-096-568A-1993	Sequence 1993, A
253	27	57.4	137	9	US-11-096-568A-5227	Sequence 5227, Ap	326	27	57.4	631	9	US-11-188-298-16078	Sequence 16078, A
254	27	57.4	149	9	US-10-530-253-17	Sequence 17, Appl1	327	27	57.4	643	11	US-11-506-454-520	Sequence 520, App
255	27	57.4	157	9	US-10-980-388-73	Sequence 73, Appl1	328	27	57.4	644	11	US-11-024-959-502	Sequence 502, App
256	27	57.4	157	11	US-11-207-078-226	Sequence 226, App	329	27	57.4	645	11	US-11-045-004-2516	Sequence 2516, Ap
257	27	57.4	168	9	US-10-454-437-196	Sequence 196, App	330	27	57.4	646	11	US-11-072-512-2502	Sequence 2502, Ap
258	27	57.4	168	9	US-11-055-822-360	Sequence 360, App	331	27	57.4	655	11	US-10-501-035-347	Sequence 347, App
259	27	57.4	168	11	US-11-058-822-10527	Sequence 10527, A	332	27	57.4	733	9	US-10-501-035-347	Sequence 26, Appl1
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261	27	57.4	182	11	US-11-170-653-49	Sequence 9124, Ap	334	27	57.4	739	11	US-11-009-063-31	Sequence 31, Appl1
262	27	57.4	182	11	US-11-087-099-9124	Sequence 2939, Ap	335	27	57.4	772	11	US-11-079-463-6679	Sequence 6679, Ap
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264	27	57.4	208	11	US-11-214-413-32	Sequence 51, Appl1	337	27	57.4	830	9	US-10-931-198-38	Sequence 38, Appl1
265	27	57.4	210	11	US-11-170-653-51	Sequence 50, Appl1	338	27	57.4	830	9	US-10-942-042-58	Sequence 58, Appl1
266	27	57.4	211	11	US-11-170-653-50	Sequence 10457, A	339	27	57.4	870	11	US-11-031-206-188	Sequence 188, App
267	27	57.4	211	11	US-11-087-099-10457	Sequence 6731, Ap	340	27	57.4	876	11	US-11-031-206-202	Sequence 202, App
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272	27	57.4	222	7	US-11-087-099-6133	Sequence 753, App	345	27	57.4	1003	11	US-11-204-755-7	Sequence 9, Appl1
273	27	57.4	229	9	US-09-978-360A-753	Sequence 522, App	346	27	57.4	1003	11	US-11-204-755-9	Sequence 11, Appl1
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275	27	57.4	229	9	US-10-973-115B-522	Sequence 522, App	348	27	57.4	1014	11	US-11-188-298-10090	Sequence 2, Appl1
276	27	57.4	229	9	US-10-137-873A-522	Sequence 522, App	349	27	57.4	1295	11	US-11-091-928-2	Sequence 1, Appl1
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278	27	57.4	229	11	US-11-290-153-522	Sequence 522, App	351	27	57.4	1532	11	US-11-212-443-62	Sequence 179, App
279	27	57.4	233	11	US-11-207-078-15	Sequence 58, Appl1	352	27	57.4	1563	9	US-11-212-443-179	Sequence 148, App
280	27	57.4	234	11	US-11-096-568A-15427	Sequence 15427, A	353	27	57.4	1663	9	US-10-055-877-143	Sequence 12160, A
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283	27	57.4	284	11	US-11-212-443-58	Sequence 58, Appl1	356	27	57.4	1978	11	US-11-045-004-38	Sequence 38, Appl1
284	27	57.4	284	11	US-11-087-099-10570	Sequence 10570, A	357	27	57.4	1981	11	US-11-045-004-38	Sequence 218, App
285	27	57.4	287	11	US-11-096-568A-2937	Sequence 2937, Ap	358	27	57.4	1985	9	US-11-052-554A-374	Sequence 374, App
286	27	57.4	288	9	US-11-525-907-44	Sequence 44, Appl1	359	27	57.4	2015	11	US-10-453-372-1142	Sequence 1142, Ap
287	27	57.4	288	9	US-11-188-298-21619	Sequence 21619, A	360	27	57.4	4913	9	US-10-453-372-1132	Sequence 1132, Ap
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289	27	57.4	295	9	US-11-058-263-50	Sequence 50, Appl1	362	27	57.4	18	11	US-11-004-399-2919	Sequence 2919, Ap
290	27	57.4	307	11	US-11-096-568A-1203	Sequence 1203, Ap	363	27	57.4	49	9	US-10-895-064-2872	Sequence 2872, Ap
291	27	57.4	321	9	US-10-508-263-52	Sequence 52, Appl1	364	27	57.4	49	11	US-11-129-741-8872	Sequence 9237, Ap
292	27	57.4	326	11	US-11-096-568A-1202	Sequence 1202, Ap	365	27	57.4	61	11	US-11-079-463-9237	Sequence 12, Appl1
293	27	57.4	327	11	US-11-072-512-2508	Sequence 2508, Ap	366	27	57.4	103	11	US-11-007-282-12	Sequence 10, Appl1
294	27	57.4	328	11	US-11-096-568A-3897	Sequence 3897, Ap	367	27	57.4	108	11	US-11-007-282-10	Sequence 544, App
295	27	57.4	334	11	US-11-096-568A-3896	Sequence 3896, Ap	368	27	57.4	129	11	US-11-176-830-544	Sequence 544, App
296	27	57.4	334	11	US-11-172-740-45	Sequence 45, Appl1	369	27	57.4	129	11	US-11-079-463-5966	Sequence 2719, Ap
297	27	57.4	339	11	US-11-096-568A-15425	Sequence 15425, A	370	27	57.4	137	11	US-11-072-512-2691	Sequence 2691, Ap
298	27	57.4	339	11	US-11-172-740-326	Sequence 326, App	371	27	57.4	140	11	US-11-072-512-2719	Sequence 2719, Ap
299	27	57.4	343	11	US-11-096-568A-6730	Sequence 6730, Ap	372	27	57.4	140	11	US-11-079-463-5966	Sequence 4212, Ap
300	27	57.4	346	11	US-11-096-568A-1201	Sequence 1201, Ap	373	27	57.4	167	11	US-11-096-568A-4212	Sequence 4212, Ap
301	27	57.4	352	11	US-11-096-568A-6729	Sequence 6729, Ap	374	27	57.4	175	11	US-11-079-463-5966	Sequence 3347, Ap
302	27	57.4	360	11	US-11-212-443-178	Sequence 178, App	375	27	57.4	180	11	US-11-188-298-291	Sequence 291, App
303	27	57.4	371	11	US-11-188-298-16916	Sequence 16916, A	376	27	57.4	181	11	US-11-072-512-3327	Sequence 3327, Ap
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305	27	57.4	388	11	US-11-096-568A-3895	Sequence 3895, A	378	27	57.4	196	9	US-10-467-657-4664	Sequence 4664, App
306	27	57.4	388	11	US-11-188-298-21843	Sequence 21843, A	379	27	57.4	212	11	US-11-096-568A-4211	Sequence 4211, Ap
307	27	57.4	401	9	US-10-878-556A-179	Sequence 179, App	380	27	57.4	214	11	US-11-096-568A-4858	Sequence 4858, Ap
308	27	57.4	437	11	US-11-087-099-5721	Sequence 5721, Ap	381	27	57.4	218	11	US-11-079-463-8535	Sequence 8535, Ap
309	27	57.4	437	11	US-11-188-298-5230	Sequence 5230, Ap	382	27	57.4				
310	27	57.4	449	9	US-10-621-234-1089	Sequence 1089, Ap	383	27	57.4				
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312	27	57.4	453	11	US-11-087-099-5622	Sequence 5622, Ap	385	27	57.4				
313	27	57.4					386	27	57.4				

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389	26	55.3	240	11	US-11-079-463-8932	Sequence 8932, Ap	462	26	55.3	536	11	US-11-124-367A-498	Sequence 498, App
390	26	55.3	241	11	US-11-096-568A-16354	Sequence 16354, A	463	26	55.3	536	11	US-11-124-367A-498	Sequence 498, App
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392	26	55.3	273	11	US-11-045-004-229	Sequence 229, App	465	26	55.3	556	9	US-10-506-454-828	Sequence 828, App
393	26	55.3	282	11	US-11-096-568A-4857	Sequence 4857, Ap	466	26	55.3	613	11	US-11-188-298-5827	Sequence 5827, Ap
394	26	55.3	293	11	US-11-087-099-513	Sequence 513, App	467	26	55.3	613	11	US-11-087-099-9717	Sequence 9717, Ap
395	26	55.3	306	11	US-11-019-711-89	Sequence 89, Appl1	468	26	55.3	664	9	US-10-542-178-1	Sequence 3, Appl1
396	26	55.3	306	11	US-11-019-711-90	Sequence 90, Appl1	469	26	55.3	721	9	US-10-542-178-1	Sequence 1, Appl1
397	26	55.3	307	10	US-11-283-522-33	Sequence 33, Appl1	470	26	55.3	748	9	US-10-455-772-300	Sequence 300, App
398	26	55.3	307	11	US-11-019-711-87	Sequence 87, Appl1	471	26	55.3	749	11	US-11-098-686-10505	Sequence 10505, A
399	26	55.3	307	11	US-11-019-711-88	Sequence 88, Appl1	472	26	55.3	754	9	US-10-455-772-294	Sequence 294, App
400	26	55.3	307	11	US-11-265-966-16	Sequence 16, Appl1	473	26	55.3	754	9	US-10-455-772-298	Sequence 298, App
401	26	55.3	330	11	US-11-072-512-2839	Sequence 2839, Ap	474	26	55.3	755	9	US-10-455-772-302	Sequence 302, App
402	26	55.3	334	11	US-11-098-686-10496	Sequence 10496, A	475	26	55.3	755	9	US-10-455-772-302	Sequence 302, App
403	26	55.3	339	11	US-11-079-463-7424	Sequence 7424, Ap	476	26	55.3	755	9	US-10-455-772-304	Sequence 304, App
404	26	55.3	348	11	US-11-096-568A-33378	Sequence 33378, A	477	26	55.3	756	9	US-10-455-772-296	Sequence 296, App
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406	26	55.3	360	9	US-10-288-733-2	Sequence 2, Appl1	479	26	55.3	777	9	US-10-645-441-3	Sequence 3, Appl1
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409	26	55.3	375	11	US-11-087-099-7023	Sequence 7023, Ap	482	26	55.3	840	9	US-10-645-441-1	Sequence 1, Appl1
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412	26	55.3	376	11	US-11-188-298-6334	Sequence 6334, Ap	485	26	55.3	841	11	US-11-050-804-2	Sequence 2, Appl1
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414	26	55.3	387	9	US-10-506-454-865	Sequence 865, App	487	26	55.3	934	9	US-10-858-730-8	Sequence 6, Appl1
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421	26	55.3	396	11	US-11-188-298-7665	Sequence 7665, Ap	494	26	55.3	1116	11	US-11-113-751-34	Sequence 34, Appl1
422	26	55.3	401	11	US-11-096-568A-33377	Sequence 33377, A	495	26	55.3	1118	11	US-11-113-751-42	Sequence 42, Appl1
423	26	55.3	409	11	US-11-096-568A-19375	Sequence 19375, A	496	26	55.3	1121	11	US-11-113-751-19	Sequence 19, Appl1
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427	26	55.3	426	11	US-11-096-568A-20044	Sequence 20044, A	500	26	55.3	1123	11	US-11-113-751-44	Sequence 44, Appl1
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430	26	55.3	432	11	US-11-096-568A-25141	Sequence 25141, A	503	26	55.3	1440	11	US-11-096-568A-34297	Sequence 34297, A
431	26	55.3	433	11	US-11-188-298-5453	Sequence 5453, Ap	504	26	55.3	1452	9	US-10-995-561-778	Sequence 778, App
432	26	55.3	435	11	US-11-087-099-11022	Sequence 11022, A	505	26	55.3	1457	9	US-10-995-561-778	Sequence 24, Appl1
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435	26	55.3	444	9	US-10-513-639-22	Sequence 22, Appl1	508	26	55.3	1866	9	US-10-511-937-186	Sequence 186, App
436	26	55.3	444	9	US-10-513-639-23	Sequence 23, Appl1	509	26	55.3	1992	11	US-11-069-834-58	Sequence 58, Appl1
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439	26	55.3	462	11	US-11-124-367A-497	Sequence 497, App	512	26	55.3	2671	8	US-10-505-928-784	Sequence 784, App
440	26	55.3	464	11	US-11-188-298-5875	Sequence 5875, Ap	513	26	55.3	2671	8	US-10-876-787-6	Sequence 6, Appl1
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442	26	55.3	466	11	US-11-188-298-2846	Sequence 2846, Ap	515	26	55.3	3960	9	US-10-995-561-771	Sequence 771, App
443	26	55.3	468	11	US-11-188-298-9547	Sequence 9547, Ap	516	26	55.3	4386	11	US-11-004-399-714	Sequence 714, App
444	26	55.3	471	11	US-11-096-568A-15988	Sequence 15988, A	517	26	55.3	4834	8	US-10-505-928-827	Sequence 827, App
445	26	55.3	472	11	US-11-188-298-1804	Sequence 1804, Ap	518	26	55.3	5335	9	US-10-995-561-777	Sequence 777, App
446	26	55.3	472	11	US-11-188-298-7009	Sequence 7009, Ap	519	26	55.3	5406	9	US-10-995-561-774	Sequence 774, App
447	26	55.3	472	11	US-11-188-298-7690	Sequence 7690, Ap	520	26	55.3	5415	9	US-10-995-561-779	Sequence 779, App
448	26	55.3	472	11	US-11-188-298-9346	Sequence 9346, Ap	521	26	55.3	5464	9	US-10-995-561-775	Sequence 775, App
449	26	55.3	472	11	US-11-188-298-15363	Sequence 15363, A	522	26	55.3	5935	9	US-10-995-561-776	Sequence 776, App
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535	25	53.2	10	11	US-11-045-024-2067	Sequence 2067, App	608	25	53.2	295	9	US-10-784-004-728	Sequence 728, App
536	25	53.2	10	11	US-11-045-024-6582	Sequence 6582, App	609	25	53.2	295	10	US-11-742-111-24	Sequence 24, App1
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539	25	53.2	11	11	US-11-045-024-5944	Sequence 5944, App	612	25	53.2	299	9	US-10-467-657-1190	Sequence 2190, App
540	25	53.2	15	11	US-11-051-481-48	Sequence 48, App1	613	25	53.2	299	11	US-11-058-924-110	Sequence 10, App1
541	25	53.2	21	9	US-10-939-890-511	Sequence 511, App	614	25	53.2	300	9	US-10-922-2328-62	Sequence 62, App1
542	25	53.2	54	11	US-11-096-568A-6075	Sequence 6075, App	615	25	53.2	303	11	US-11-058-924-5	Sequence 5, App1
543	25	53.2	64	9	US-10-467-657-3398	Sequence 3398, App	616	25	53.2	303	11	US-11-172-740-1720	Sequence 1720, App
544	25	53.2	66	11	US-11-051-481-45	Sequence 45, App1	617	25	53.2	303	11	US-11-045-004-1069	Sequence 1069, App
545	25	53.2	79	11	US-11-264-096-406	Sequence 406, App	618	25	53.2	307	9	US-10-793-626-1122	Sequence 2122, App
546	25	53.2	79	11	US-11-264-096-407	Sequence 407, App	619	25	53.2	311	9	US-10-467-657-6644	Sequence 10813, App
547	25	53.2	82	11	US-11-096-568A-1519	Sequence 1519, App	620	25	53.2	316	11	US-11-058-924-10013	Sequence 2, App1
548	25	53.2	86	11	US-11-051-481-44	Sequence 44, App1	621	25	53.2	316	11	US-11-096-568A-2807	Sequence 2807, App
549	25	53.2	88	11	US-11-096-568A-21029	Sequence 21029, A	622	25	53.2	316	11	US-11-096-568A-2810	Sequence 2810, App
550	25	53.2	102	11	US-11-264-096-1133	Sequence 1133, App	623	25	53.2	317	11	US-11-094-519A-38	Sequence 15, App1
551	25	53.2	105	11	US-11-073-420-31	Sequence 31, App1	624	25	53.2	322	11	US-11-067-121-15	Sequence 58, App1
552	25	53.2	108	11	US-11-096-568A-3660	Sequence 3660, App	625	25	53.2	322	9	US-10-921-793-58	Sequence 58, App1
553	25	53.2	113	11	US-11-072-512-2416	Sequence 2416, App	626	25	53.2	323	9	US-10-931-198-58	Sequence 58, App1
554	25	53.2	113	11	US-11-176-830-574	Sequence 574, App	627	25	53.2	323	9	US-10-942-042-58	Sequence 58, App1
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558	25	53.2	118	11	US-11-079-463-5839	Sequence 5839, App	631	25	53.2	326	11	US-11-228-875-75	Sequence 75, App1
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562	25	53.2	125	11	US-11-194-246-395	Sequence 395, App	635	25	53.2	330	11	US-11-096-568A-7841	Sequence 7841, App
563	25	53.2	127	11	US-11-197-380-2	Sequence 2, App1	636	25	53.2	330	11	US-11-096-568A-28576	Sequence 28576, A
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565	25	53.2	128	9	US-10-821-234-11229	Sequence 1229, App	638	25	53.2	334	11	US-11-165-211-53	Sequence 53, App1
566	25	53.2	135	11	US-11-045-004-1028	Sequence 1028, App	639	25	53.2	334	11	US-11-165-226-63	Sequence 63, App1
567	25	53.2	139	9	US-10-467-657-7880	Sequence 7880, App	640	25	53.2	334	11	US-11-188-298-9106	Sequence 9106, App
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570	25	53.2	150	11	US-11-188-298-11815	Sequence 11815, A	643	25	53.2	340	11	US-11-188-298-7840	Sequence 7840, App
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573	25	53.2	158	9	US-10-530-253-26	Sequence 26, App1	646	25	53.2	344	11	US-11-079-463-6695	Sequence 6695, App
574	25	53.2	166	11	US-11-094-519A-47	Sequence 47, App1	647	25	53.2	345	11	US-11-124-368A-285	Sequence 285, App
575	25	53.2	168	9	US-10-927-641-69	Sequence 69, App1	648	25	53.2	345	11	US-11-188-298-1265	Sequence 1265, App
576	25	53.2	170	11	US-11-232-406A-26	Sequence 26, App1	649	25	53.2	347	9	US-10-516-635-6	Sequence 6, App1
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581	25	53.2	201	11	US-11-096-568A-33823	Sequence 33823, A	654	25	53.2	349	9	US-10-942-042-56	Sequence 56, App1
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583	25	53.2	209	11	US-11-045-004-568	Sequence 568, App	656	25	53.2	349	11	US-11-067-121-14	Sequence 14, App1
584	25	53.2	216	11	US-11-096-568A-31555	Sequence 31555, A	657	25	53.2	352	11	US-11-172-740-1103	Sequence 1103, App
585	25	53.2	223	11	US-11-172-740-1412	Sequence 1412, App	658	25	53.2	355	11	US-11-098-686-10134	Sequence 10134, A
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587	25	53.2	237	9	US-10-527-771-12	Sequence 12, App1	660	25	53.2	362	11	US-11-098-686-11051	Sequence 11051, A
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590	25	53.2	243	11	US-11-079-463-9859	Sequence 9859, App	663	25	53.2	369	9	US-10-467-657-2936	Sequence 2936, App
591	25	53.2	247	11	US-11-113-424-76	Sequence 76, App1	664	25	53.2	369	11	US-11-087-099-653	Sequence 653, App
592	25	53.2	247	11	US-11-172-740-824	Sequence 824, App	665	25	53.2	369	11	US-11-087-099-4196	Sequence 4196, App
593	25	53.2	250	11	US-11-054-515-1574	Sequence 1574, App	666	25	53.2	369	11	US-11-087-099-7687	Sequence 7687, App
594	25	53.2	250	11	US-11-266-444-1574	Sequence 1574, App	667	25	53.2	370	11	US-11-087-099-10170	Sequence 10170, A
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604	25	53.2	283	11	US-11-096-568A-5964	Sequence 5964, App	677	25	53.2	375	11	US-11-087-099-12120	Sequence 12120, A
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732	25	53.2	561	11	US-11-188-298-7184	Sequence 7184, Ap	805	25	53.2	1542	9	US-10-453-372-266	Sequence 266, Appl1
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745	25	53.2	712	11	US-11-037-243-69	Sequence 69, Appl1	818	25	53.2	1966	9	US-10-480-330-16	Sequence 16, Appl1
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827	25	53.2	2409	11	US-11-183-261-38	Sequence 38, App1	900	24	51.1	132	9	US-10-498-026-39	Sequence 39, App1
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831	25	53.2	2628	9	US-10-455-772-502	Sequence 502, App	904	24	51.1	142	11	US-11-045-208-39	Sequence 39, App1
832	25	53.2	2715	11	US-11-096-051-2	Sequence 2, App1	905	24	51.1	143	11	US-11-264-096-1762	Sequence 1762, Ap
833	25	53.2	2715	11	US-11-113-424-51	Sequence 51, App1	906	24	51.1	143	11	US-11-264-096-1762	Sequence 1762, Ap
834	25	53.2	2721	11	US-10-455-772-522	Sequence 522, App	907	24	51.1	148	9	US-10-821-234-869	Sequence 869, App
835	25	53.2	2721	11	US-11-096-051-10	Sequence 10, App1	908	24	51.1	151	9	US-10-667-295-220	Sequence 220, App
836	25	53.2	2725	9	US-10-455-772-486	Sequence 486, App	909	24	51.1	151	11	US-11-087-099-5350	Sequence 5350, Ap
837	25	53.2	2725	9	US-10-455-772-526	Sequence 526, App	910	24	51.1	152	11	US-11-069-642-7	Sequence 7, App1
838	25	53.2	2725	9	US-10-455-772-544	Sequence 544, App	911	24	51.1	152	11	US-10-530-253-23	Sequence 23, App1
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842	25	53.2	2725	9	US-10-455-772-552	Sequence 552, App	915	24	51.1	158	9	US-10-467-657-3592	Sequence 3592, Ap
843	25	53.2	2725	11	US-11-096-051-8	Sequence 8, App1	916	24	51.1	158	9	US-10-467-657-3592	Sequence 3592, Ap
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845	25	53.2	2801	11	US-11-124-367A-433	Sequence 433, App	918	24	51.1	158	9	US-10-467-657-3592	Sequence 3592, Ap
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847	25	53.2	2896	11	US-11-124-367A-434	Sequence 434, App	920	24	51.1	165	11	US-11-175-690-99	Sequence 99, App1
848	25	53.2	3256	11	US-11-124-368A-304	Sequence 304, App	921	24	51.1	165	11	US-11-175-690-517	Sequence 517, App
849	25	53.2	3256	11	US-11-124-367A-432	Sequence 432, App	922	24	51.1	165	11	US-11-132-722-45	Sequence 45, App1
850	25	53.2	3343	11	US-11-122-396-7	Sequence 7, App1	923	24	51.1	166	11	US-11-176-830-186	Sequence 186, App
851	25	53.2	3396	8	US-10-505-928-449	Sequence 449, App	924	24	51.1	166	11	US-11-176-830-186	Sequence 186, App
852	25	53.2	3396	11	US-11-183-261-39	Sequence 39, App1	925	24	51.1	166	11	US-11-198-765-83	Sequence 83, App1
853	25	53.2	3623	9	US-10-995-561-593	Sequence 593, App	926	24	51.1	167	11	US-11-175-690-103	Sequence 103, App
854	25	53.2	4644	9	US-10-784-004-704	Sequence 704, App	927	24	51.1	168	11	US-11-135-603-12	Sequence 12, App1
855	25	53.2	4644	9	US-10-784-004-383	Sequence 383, App	928	24	51.1	170	9	US-10-508-263-6	Sequence 6, App1
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859	24.5	52.1	270	9	US-10-495-597-4	Sequence 4, App1	932	24	51.1	172	11	US-11-172-740-601	Sequence 601, App
860	24	51.1	9	9	US-10-530-061-102	Sequence 102, App	933	24	51.1	180	11	US-11-172-740-601	Sequence 1704, A
861	24	51.1	9	9	US-10-530-061-836	Sequence 836, App	934	24	51.1	181	11	US-11-188-298-19399	Sequence 19399, A
862	24	51.1	9	9	US-10-530-061-837	Sequence 837, App	935	24	51.1	181	11	US-11-079-463-5704	Sequence 5704, Ap
863	24	51.1	11	9	US-10-530-061-494	Sequence 494, App	936	24	51.1	181	11	US-11-264-096-910	Sequence 910, App
864	24	51.1	14	11	US-11-004-399-1532	Sequence 1532, Ap	937	24	51.1	184	11	US-11-096-568A-28413	Sequence 28413, A
865	24	51.1	15	9	US-10-530-061-1707	Sequence 1707, Ap	938	24	51.1	186	11	US-11-045-004-339	Sequence 339, App
866	24	51.1	19	9	US-11-004-399-991	Sequence 991, App	939	24	51.1	189	11	US-11-087-099-9329	Sequence 929, App
867	24	51.1	31	11	US-11-004-399-767	Sequence 767, App	940	24	51.1	189	11	US-11-147-492-22	Sequence 6, App1
868	24	51.1	35	9	US-10-821-234-1704	Sequence 1704, Ap	941	24	51.1	190	11	US-11-147-492-22	Sequence 22, App1
869	24	51.1	42	9	US-10-957-887B-100	Sequence 100, App	942	24	51.1	192	11	US-11-069-601-13	Sequence 13, App1
870	24	51.1	48	11	US-11-000-463-341	Sequence 341, App	943	24	51.1	194	8	US-11-188-298-9581	Sequence 9581, Ap
871	24	51.1	65	11	US-11-000-463-813	Sequence 813, App	944	24	51.1	196	11	US-11-188-298-14820	Sequence 14820, A
872	24	51.1	65	11	US-11-096-568A-27328	Sequence 27328, A	945	24	51.1	196	11	US-11-264-096-1622	Sequence 1622, Ap
873	24	51.1	67	11	US-11-188-298-2953	Sequence 2953, Ap	946	24	51.1	199	8	US-10-511-937-2440	Sequence 2440, Ap
874	24	51.1	69	11	US-11-096-568A-2425	Sequence 2425, Ap	947	24	51.1	199	9	US-10-647-072-2	Sequence 2, App1
875	24	51.1	74	9	US-10-467-657-3524	Sequence 3524, Ap	948	24	51.1	199	11	US-11-264-096-113	Sequence 113, App
876	24	51.1	75	11	US-11-096-568A-3452	Sequence 3452, Ap	949	24	51.1	199	11	US-11-264-096-114	Sequence 114, App
877	24	51.1	80	11	US-11-096-568A-2424	Sequence 2424, Ap	950	24	51.1	204	11	US-11-096-568A-20799	Sequence 20799, A
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879	24	51.1	81	11	US-11-051-720-1309	Sequence 1309, Ap	952	24	51.1	204	11	US-11-209-188-16	Sequence 16, App1
880	24	51.1	82	9	US-10-506-454-201	Sequence 201, App	953	24	51.1	206	11	US-11-096-568A-26047	Sequence 26047, A
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882	24	51.1	90	11	US-11-205-109-12	Sequence 12, App1	955	24	51.1	213	11	US-11-188-298-16073	Sequence 16073, A
883	24	51.1	101	9	US-10-469-469-11	Sequence 11, App1	956	24	51.1	213	11	US-11-188-298-19337	Sequence 19337, A
884	24	51.1	102	11	US-11-072-512-3253	Sequence 3253, Ap	957	24	51.1	213	11	US-11-188-298-20640	Sequence 20640, A
885	24	51.1	105	11	US-11-096-568A-30678	Sequence 30678, A	958	24	51.1	215	11	US-11-045-004-2686	Sequence 2686, Ap
886	24	51.1	106	11	US-11-096-568A-6550	Sequence 6550, Ap	959	24	51.1	218	11	US-11-098-686-10245	Sequence 10245, A
887	24	51.1	113	11	US-11-072-512-3055	Sequence 3055, Ap	960	24	51.1	219	11	US-11-096-568A-6353	Sequence 6353, Ap
888	24	51.1	113	11	US-11-087-099-6180	Sequence 6180, Ap	961	24	51.1	219	11	US-11-096-568A-22484	Sequence 22484, A
889	24	51.1	114	9	US-10-241-975-52	Sequence 52, App1	962	24	51.1	221	11	US-11-051-720-1776	Sequence 1776, Ap
890	24	51.1	114	11	US-11-051-720-1425	Sequence 1425, Ap	963	24	51.1	221	11	US-11-043-806-574	Sequence 574, App
891	24	51.1	116	11	US-11-188-298-5279	Sequence 5279, Ap	964	24	51.1	225	9	US-10-469-469-7	Sequence 7, App1
892	24	51.1	118	11	US-11-116-144-156	Sequence 156, App	965	24	51.1	225	11	US-11-156-516-38	Sequence 38, App1
893	24	51.1	118	11	US-11-220-372-156	Sequence 156, App	966	24	51.1	226	11	US-11-096-568A-26045	Sequence 26045, A
894	24	51.1	127	11	US-11-186-284-171	Sequence 171, App	967	24	51.1	230	11	US-11-098-686-10984	Sequence 10984, A
895	24	51.1	129	11	US-11-116-144-145	Sequence 145, App	968	24	51.1	232	11	US-11-072-512-3602	Sequence 3602, Ap
896	24	51.1	129	11	US-11-220-372-145	Sequence 145, App	969	24	51.1	232	11	US-11-052-554A-332	Sequence 332, App
897	24	51.1	131	11	US-11-188-298-5337	Sequence 5337, Ap	970	24	51.1	236	11	US-11-123-241-56	Sequence 56, App1

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971 24 51.1 238 11 US-11-188-298-7776 Sequence 7776, Ap
972 24 51.1 239 9 US-11-045-1004-139 Sequence 139, App
973 24 51.1 247 11 US-10-632-150-36 Sequence 36, Appl
974 24 51.1 247 10 US-11-106-014-36 Sequence 36, Appl
975 24 51.1 247 11 US-11-073-457-36 Sequence 36, Appl
976 24 51.1 247 11 US-11-073-460-36 Sequence 36, Appl
977 24 51.1 250 11 US-11-087-099-3334 Sequence 3334, Ap
978 24 51.1 250 11 US-11-096-568A-1943 Sequence 1943, Ap
979 24 51.1 252 11 US-11-229-769-214 Sequence 214, App
980 24 51.1 253 11 US-11-054-515-1200 Sequence 1200, Ap
981 24 51.1 253 11 US-11-266-444-1200 Sequence 1200, Ap
982 24 51.1 254 11 US-11-096-568A-28412 Sequence 28412, A
983 24 51.1 258 11 US-11-188-298-11673 Sequence 11673, A
984 24 51.1 259 9 US-10-986-405-312 Sequence 312, App
985 24 51.1 263 11 US-11-072-512-3517 Sequence 3517, Ap
986 24 51.1 267 9 US-10-793-626-1508 Sequence 1508, Ap
987 24 51.1 267 11 US-11-140-024-5 Sequence 5, Appl
988 24 51.1 267 11 US-11-096-568A-17567 Sequence 17567, A
989 24 51.1 269 11 US-11-188-298-16681 Sequence 16681, A
990 24 51.1 270 9 US-10-467-657-3022 Sequence 3022, Ap
991 24 51.1 270 9 US-10-467-657-5806 Sequence 5806, Ap
992 24 51.1 273 9 US-10-793-626-166 Sequence 166, App
993 24 51.1 276 11 US-11-096-568A-28559 Sequence 28559, A
994 24 51.1 279 11 US-11-096-568A-33858 Sequence 33858, A
995 24 51.1 281 11 US-11-087-099-7170 Sequence 7170, Ap
996 24 51.1 282 11 US-11-096-568A-15872 Sequence 15872, A
997 24 51.1 284 9 US-10-469-469-4 Sequence 4, Appl
998 24 51.1 284 11 US-11-188-298-579 Sequence 579, App
999 24 51.1 286 9 US-10-455-772-430 Sequence 430, App
1000 24 51.1 290 11 US-11-207-626A-17 Sequence 17, Appl
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ALIGNMENTS

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RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13
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Query Match 100.0%; Score 47; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.022;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy 1 QLCLELQTT 9
Db 14 QLCLELQTT 22
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RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1
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; GENERAL INFORMATION:
; APPLICANT: Healthbarks Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; PRIOR FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3
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Query Match 100.0%; Score 47; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.023;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy 1 QLCLELQTT 9
Db 21 QLCLELQTT 29
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RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-1
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Query Match 100.0%; Score 47; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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Qy 1 QLCLELQTT 9
Db 14 QLCLELQTT 22
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RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
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PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 100.0%; Score 47; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 14 QCTELQTT 22

RESULT 5
US-10-530-253-5
Sequence 5, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 47; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 14 QCTELQTT 22

RESULT 6
US-10-530-253-7
Sequence 7, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
LENGTH: 248

TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 100.0%; Score 47; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 111 QCTELQTT 119

RESULT 7
US-10-530-253-9
Sequence 9, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 47; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELQTT 9
Db 111 QCTELQTT 119

RESULT 8
US-10-530-253-11
Sequence 11, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 11
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 47; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELOT 9
Db 111 QCTELOT 119

RESULT 9

US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; TITLE OF INVENTION: RECOMBINANT REPLICON
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; CURRENT FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 47; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.037;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 QCTELOT 9
Db 119 QCTELOT 127

RESULT 10

US-11-098-662-8
; Sequence 8, Application US/11098662
; Publication No. US2005024423A1
; GENERAL INFORMATION:
; APPLICANT: Klucher, Kevin M.
; APPLICANT: Kindvogel, Wayne R.
; APPLICANT: Sivakumar, Pallavur V.
; APPLICANT: Henderson, Katherine E.
; TITLE OF INVENTION: METHODS FOR TREATING VIRAL INFECTION
; TITLE OF INVENTION: USING IL-28 AND IL-29 CYSTEINE MUTANTS
; FILE REFERENCE: 04-05
; CURRENT APPLICATION NUMBER: US/11/098,662
; CURRENT FILING DATE: 2005-04-04
; NUMBER OF SEQ ID NOS: 136
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Mus musculus
US-11-098-662-8

Query Match 72.3%; Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 QCTELOT 9
Db 139 QCTELOT 147

RESULT 11
US-11-098-662-10

; Sequence 10, Application US/11098662
; Publication No. US2005024423A1
; GENERAL INFORMATION:
; APPLICANT: Klucher, Kevin M.
; APPLICANT: Kindvogel, Wayne R.
; APPLICANT: Sivakumar, Pallavur V.
; APPLICANT: Henderson, Katherine E.
; TITLE OF INVENTION: METHODS FOR TREATING VIRAL INFECTION
; TITLE OF INVENTION: USING IL-28 AND IL-29 CYSTEINE MUTANTS
; FILE REFERENCE: 04-05
; CURRENT APPLICATION NUMBER: US/11/098,662
; CURRENT FILING DATE: 2005-04-04
; NUMBER OF SEQ ID NOS: 136
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Mus musculus
US-11-098-662-10

Query Match 72.3%; Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 QCTELOT 9
Db 139 QCTELOT 147

RESULT 12

US-11-193-955-8
; Sequence 8, Application US/11193955
; Publication No. US20060024269A1
; GENERAL INFORMATION:
; APPLICANT: DOYLE, SEAN
; APPLICANT: KLUCHER, KEVIN M.
; APPLICANT: SIVAKUMAR, PALLAVUR V.
; APPLICANT: KINDSVOGEL, WAYNE R.
; APPLICANT: CHAN, CHUNG
; TITLE OF INVENTION: USE OF IL-28 AND IL-29 TO TREAT CANCER
; TITLE OF INVENTION: AND AUTOIMMUNE DISORDERS
; FILE REFERENCE: 04-08
; CURRENT APPLICATION NUMBER: US/11/193,955
; CURRENT FILING DATE: 2005-07-29
; NUMBER OF SEQ ID NOS: 161
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Mus musculus
; FEATURE:
; NAME/KEY: SIGNAL
; LOCATION: (1)...(28)
US-11-193-955-8

Query Match 72.3%; Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

Qy 1 QCTELOT 9
Db 139 QCTELOT 147

RESULT 13

US-11-193-955-10
; Sequence 10, Application US/11193955
; Publication No. US20060024269A1
; GENERAL INFORMATION:
; APPLICANT: DOYLE, SEAN
; APPLICANT: KLUCHER, KEVIN M.
; APPLICANT: SIVAKUMAR, PALLAVUR V.
; APPLICANT: KINDSVOGEL, WAYNE R.

APPLICANT: CHAN, CHUNG
TITLE OF INVENTION: USE OF IL-28 AND IL-29 TO TREAT CANCER
FILE REFERENCE: 04-08
CURRENT APPLICATION NUMBER: US/11/193,955
CURRENT FILING DATE: 2005-07-29
NUMBER OF SEQ ID NOS: 161
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 10
LENGTH: 202
TYPE: PRT
ORGANISM: Mus musculus
FEATURE:
NAME/KEY: SIGNAL
LOCATION: (1)...(28)
US-11-193-955-10

Query Match 72.3% Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

OY 1 QCTELQTT 9
Db 139 QCTQLOQT 147

RESULT 14
US-11-266-626-9
Sequence 9, Application US/11266626
Publication No. US20060063233A1
GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
APPLICANT: Fox, Brian A.
APPLICANT: Klucher, Kevin M.
APPLICANT: Taft, David W.
APPLICANT: Kindsvogel, Wayne R.
TITLE OF INVENTION: CYTOKINE PROTEIN FAMILY
FILE REFERENCE: 01-17
CURRENT APPLICATION NUMBER: US/11/266,626
CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: US/10/127,816
PRIOR FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: US 60/285,408
PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: US 60/286,482
PRIOR FILING DATE: 2001-04-25
PRIOR APPLICATION NUMBER: US 60/341,050
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/341,105
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 09/895,834
PRIOR FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/285,424
PRIOR FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 59
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 9
LENGTH: 202
TYPE: PRT
ORGANISM: Homo sapiens
US-11-266-626-9

Query Match 72.3% Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

OY 1 QCTELQTT 9
Db 139 QCTQLOQT 147

RESULT 15
US-11-266-626-11

Sequence 11, Application US/11266626
Publication No. US20060063233A1
GENERAL INFORMATION:
APPLICANT: Sheppard, Paul O.
APPLICANT: Fox, Brian A.
APPLICANT: Klucher, Kevin M.
APPLICANT: Taft, David W.
APPLICANT: Kindsvogel, Wayne R.
TITLE OF INVENTION: CYTOKINE PROTEIN FAMILY
FILE REFERENCE: 01-17
CURRENT APPLICATION NUMBER: US/11/266,626
CURRENT FILING DATE: 2005-11-03
PRIOR APPLICATION NUMBER: US/10/127,816
PRIOR FILING DATE: 2002-04-19
PRIOR APPLICATION NUMBER: US 60/285,408
PRIOR FILING DATE: 2001-04-20
PRIOR APPLICATION NUMBER: US 60/286,482
PRIOR FILING DATE: 2001-04-25
PRIOR APPLICATION NUMBER: US 60/341,050
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 60/341,105
PRIOR FILING DATE: 2001-10-22
PRIOR APPLICATION NUMBER: US 09/895,834
PRIOR FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/285,424
PRIOR FILING DATE: 2001-04-20
NUMBER OF SEQ ID NOS: 59
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 11
LENGTH: 202
TYPE: PRT
ORGANISM: Homo sapiens
US-11-266-626-11

Query Match 72.3% Score 34; DB 11; Length 202;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

OY 1 QCTELQTT 9
Db 139 QCTQLOQT 147

RESULT 16
US-10-530-253-15
Sequence 15, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Cassetti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/1004137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 15
LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 18
US-10-530-253-15

Query Match 70.2% Score 33; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 15;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 2 LCTELQTT 9

Db 17 LCTELNTS 24

RESULT 17
US-10-530-253-20
; Sequence 20, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/1100137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 45
US-10-530-253-20

Query Match 70.2%; Score 33; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 15;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 17 LCTELNTS 24

RESULT 18
US-11-079-463-6325
; Sequence 6325, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FR
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6325
; LENGTH: 254
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-6325

Query Match 70.2%; Score 33; DB 11; Length 254;
Best Local Similarity 75.0%; Pred. No. 24;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 78 LCTELQTT 85

RESULT 19
US-11-087-099-6055
; Sequence 6055, Application US/11087099
; Publication No. US20060041961A1

; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 6055
; LENGTH: 420
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)-(420)
; OTHER INFORMATION: unsure at all Xaa locations
US-11-087-099-6055

Query Match 70.2%; Score 33; DB 11; Length 420;
Best Local Similarity 85.7%; Pred. No. 39;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 3 CTELOTT 9
Db 308 CTELOTT 314

RESULT 20
US-11-072-512-2475
; Sequence 2475, Application US/11072512
; Publication No. US20060029945A1
; GENERAL INFORMATION:
; APPLICANT: ISOGAI, TAKAO
; APPLICANT: SUGIYAMA, TOMOYASU
; APPLICANT: OTSUKI, TETSUJI
; APPLICANT: WAKAMATSU, AI
; APPLICANT: SATO, HIROYUKI
; APPLICANT: ISHII, SHIZUKO
; APPLICANT: YAMAMOTO, JUN-ICHI
; APPLICANT: ISONO, YUUKO
; APPLICANT: HIO, YUKI
; APPLICANT: OTSUKA, KAORU
; APPLICANT: NAGAI, KEIICHI
; APPLICANT: IRIE, RYOTARO
; APPLICANT: TAMECHIKI, ICHIRO
; APPLICANT: SEKI, NAOHITO
; APPLICANT: YOSHIKAWA, TSUTOMU
; APPLICANT: OTSUKA, MOTOYUKI
; APPLICANT: NAGAHARI, KENJI
; APPLICANT: MASUHO, YASUHIKO
; TITLE OF INVENTION: Novel full length cDNA
; FILE REFERENCE: 084335-0191
; CURRENT APPLICATION NUMBER: US/11/072,512
; CURRENT FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US 60/350,978
; PRIOR FILING DATE: 2002-01-25
; PRIOR FILING DATE: 2001-11-05
; PRIOR APPLICATION NUMBER: JP 2001-379298
; NUMBER OF SEQ ID NOS: 4096
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2475
; LENGTH: 1098
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-072-512-2475

Query Match 70.2%; Score 33; DB 11; Length 1098;
Best Local Similarity 85.7%; Pred. No. 96;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 QCTELOQ 7
Db 285 QCTELOQ 291

RESULT 21

US-10-467-7248
Sequence 7248, Application US/10467657
Publication No. US20050260581A1
GENERAL INFORMATION:
APPLICANT: CHIRON SPA
APPLICANT: FONTANA Maria Rita
APPLICANT: PIZZA Mariagrazia
APPLICANT: MASIGNANI Vega
TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqMan9, version 1.04
SEQ ID NO 7248
LENGTH: 43
TYPE: PRT
ORGANISM: Neisseria gonorrhoeae
US-10-467-657-7248

Query Match 66.0%; Score 31; DB 9; Length 43;
Best Local Similarity 71.4%; Pred. No. 11;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 3 CTELOTT 9
|||||

DB 13 CTELEST 19

RESULT 22

US-10-519-390-15
Sequence 15, Application US/10519390
Publication No. US20060008872A1
GENERAL INFORMATION:
APPLICANT: MEDENGEN Inc.
APPLICANT: CHUNG, Yong-Hoon
APPLICANT: LEE, Hak-sup
APPLICANT: YI, Ki-Man
APPLICANT: KIM, Jae-Youn
APPLICANT: HEO, Youn-Hwa
TITLE OF INVENTION: A method of improving efficacy of biological response-modifying
TITLE OF INVENTION: proteins and the example mutants
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/519,390
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: KR10-2003-0051846
PRIOR FILING DATE: 2003-07-26
NUMBER OF SEQ ID NOS: 65
SOFTWARE: Koparentin 1.71
SEQ ID NO 15
LENGTH: 129
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: IL-4: 33rd, 45th, 55th, 73rd, 82nd or 112nd phe is replaced by
OTHER INFORMATION: Val.
US-10-519-390-15

Query Match 66.0%; Score 31; DB 9; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELOTT 9
|||||

DB 23 LCTELTVT 30

RESULT 23

US-11-176-830-207

Sequence 207, Application US/11176830
Publication No. US2006020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 207
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: Genbank AAA59149
DATABASE ENTRY DATE: 1995-01-06
US-11-176-830-207

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELOTT 9
|||||

DB 23 LCTELTVT 30

RESULT 24

US-11-176-830-546
Sequence 546, Application US/11176830
Publication No. US2006020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
APPLICANT: Vega, Manuel
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 546
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-546

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELOTT 9
|||||

Db 23 LCTELTVT 30

```
RESULT 25
US-11-176-830-547
; Sequence 547, Application US/11176830
; Publication No. US2006020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 547
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-547
```

```
Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

```
RESULT 26
US-11-176-830-548
; Sequence 548, Application US/11176830
; Publication No. US2006020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 548
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-548
```

```
Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

Qy 2 LCTELQTT 9

Db 23 LCTELTVT 30

```
RESULT 27
US-11-176-830-549
; Sequence 549, Application US/11176830
; Publication No. US2006020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 549
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-549
```

```
Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

```
RESULT 28
US-11-176-830-550
; Sequence 550, Application US/11176830
; Publication No. US2006020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 550
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-550
```

```
Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

Qy 2 LCTELQTT 9
| | | | |
Db 23 LCTELTVT 30

RESULT 29
US-11-176-830-551
; Sequence 551, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dricianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT FILING DATE: 2005-07-06
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 551
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-551

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
| | | | |
Db 23 LCTELTVT 30

RESULT 30
US-11-176-830-552
; Sequence 552, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dricianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT FILING DATE: 2005-07-06
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 552
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-552

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
| | | | |
Db 23 LCTELTVT 30

RESULT 31
US-11-176-830-553
; Sequence 553, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dricianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT FILING DATE: 2005-07-06
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 553
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-553

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
| | | | |
Db 23 LCTELTVT 30

RESULT 32
US-11-176-830-554
; Sequence 554, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dricianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT FILING DATE: 2005-07-06
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 554
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-554

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 23 LCTELTWT 30

RESULT 33
US-11-176-830-555
; Sequence 555, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 555
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-555

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 23 LCTELTWT 30

RESULT 34
US-11-176-830-556
; Sequence 556, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 556
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-556

Query Match 66.0%; Score 31; DB 11; Length 129;

Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 23 LCTELTWT 30

RESULT 35
US-11-176-830-557
; Sequence 557, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 557
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-557

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
Db 23 LCTELTWT 30

RESULT 36
US-11-176-830-558
; Sequence 558, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 558
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-558

Query Match 66.0%; Score 31; DB 11; Length 129;

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 37

US-11-176-830-559
; Sequence 559, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Diltant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 559
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-559

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 38

US-11-176-830-560
; Sequence 560, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Diltant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 560
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-560

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 39

US-11-176-830-561
; Sequence 561, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Diltant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 561
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-561

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 40

US-11-176-830-562
; Sequence 562, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Diltant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 562
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-562

US-11-176-830-562

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

RESULT 41

US-11-176-830-563

; Sequence 563, Application US/11176830
; Publication No. US2006020116A1

; GENERAL INFORMATION:

; APPLICANT: Gantier, Rene

; APPLICANT: Guyon, Thierry

; APPLICANT: Dittanci, Lila

; APPLICANT: Vega, Manuel

; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc

; FILE REFERENCE: 17109-012002 (922B)

; CURRENT APPLICATION NUMBER: US/11/176, 830

; CURRENT FILING DATE: 2005-07-06

; PRIOR APPLICATION NUMBER: 10/658, 834

; PRIOR FILING DATE: 2003-09-08

; PRIOR APPLICATION NUMBER: 60/457, 135

; PRIOR FILING DATE: 2003-03-21

; PRIOR APPLICATION NUMBER: 60/409, 898

; PRIOR FILING DATE: 2002-09-09

; NUMBER OF SEQ ID NOS: 1306

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 563

; LENGTH: 129

; TYPE: PRT

; ORGANISM: Homo sapiens

US-11-176-830-563

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

RESULT 42

US-11-176-830-564

; Sequence 564, Application US/11176830
; Publication No. US2006020116A1

; GENERAL INFORMATION:

; APPLICANT: Gantier, Rene

; APPLICANT: Guyon, Thierry

; APPLICANT: Dittanci, Lila

; APPLICANT: Vega, Manuel

; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc

; FILE REFERENCE: 17109-012002 (922B)

; CURRENT APPLICATION NUMBER: US/11/176, 830

; CURRENT FILING DATE: 2005-07-06

; PRIOR APPLICATION NUMBER: 10/658, 834

; PRIOR FILING DATE: 2003-09-08

; PRIOR APPLICATION NUMBER: 60/457, 135

; PRIOR FILING DATE: 2003-03-21

; PRIOR APPLICATION NUMBER: 60/409, 898

; PRIOR FILING DATE: 2002-09-09

; NUMBER OF SEQ ID NOS: 1306

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 564

; LENGTH: 129

; TYPE: PRT

; ORGANISM: Homo sapiens
US-11-176-830-564

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

RESULT 43

US-11-176-830-565

; Sequence 565, Application US/11176830
; Publication No. US2006020116A1

; GENERAL INFORMATION:

; APPLICANT: Gantier, Rene

; APPLICANT: Guyon, Thierry

; APPLICANT: Dittanci, Lila

; APPLICANT: Vega, Manuel

; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc

; FILE REFERENCE: 17109-012002 (922B)

; CURRENT APPLICATION NUMBER: US/11/176, 830

; CURRENT FILING DATE: 2005-07-06

; PRIOR APPLICATION NUMBER: 10/658, 834

; PRIOR FILING DATE: 2003-09-08

; PRIOR APPLICATION NUMBER: 60/457, 135

; PRIOR FILING DATE: 2003-03-21

; PRIOR APPLICATION NUMBER: 60/409, 898

; PRIOR FILING DATE: 2002-09-09

; NUMBER OF SEQ ID NOS: 1306

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 565

; LENGTH: 129

; TYPE: PRT

; ORGANISM: Homo sapiens

US-11-176-830-565

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 2 LCTELQTT 9
Db 23 LCTELTVT 30

RESULT 44

US-11-176-830-566

; Sequence 566, Application US/11176830
; Publication No. US2006020116A1

; GENERAL INFORMATION:

; APPLICANT: Gantier, Rene

; APPLICANT: Guyon, Thierry

; APPLICANT: Dittanci, Lila

; APPLICANT: Vega, Manuel

; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc

; FILE REFERENCE: 17109-012002 (922B)

; CURRENT APPLICATION NUMBER: US/11/176, 830

; CURRENT FILING DATE: 2005-07-06

; PRIOR APPLICATION NUMBER: 10/658, 834

; PRIOR FILING DATE: 2003-09-08

; PRIOR APPLICATION NUMBER: 60/457, 135

; PRIOR FILING DATE: 2003-03-21

; PRIOR APPLICATION NUMBER: 60/409, 898

; PRIOR FILING DATE: 2002-09-09

; NUMBER OF SEQ ID NOS: 1306

; SOFTWARE: FastSeq for Windows Version 4.0

; SEQ ID NO 566

; LENGTH: 129

TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-566

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 45

US-11-176-830-567
Sequence 567, Application US/11176830
Publication No. US20060020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Drltanti, Lila
APPLICANT: Vega, Manuel
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 567
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-567

Query Match 66.0%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 23 LCTELTVT 30

RESULT 46

US-10-469-561-25
Sequence 25, Application US/10469561
Publication No. US20050260216A1
GENERAL INFORMATION:
APPLICANT: Claire Ashman
APPLICANT: James Scott Crowe
APPLICANT: Jonathan Henry Ellis
APPLICANT: Alan Peter Lewis
TITLE OF INVENTION: VACCINE
FILE REFERENCE: PG4355USW
CURRENT APPLICATION NUMBER: US/10/469,561
CURRENT FILING DATE: 2003-08-29
NUMBER OF SEQ ID NOS: 25
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 25
LENGTH: 150
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimeric IL4 for human use
US-10-469-561-25

Query Match 66.0%; Score 31; DB 9; Length 150;
Best Local Similarity 75.0%; Pred. No. 36;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 48 LCTELTVT 55

RESULT 47

US-10-511-937-2476
Sequence 2476, Application US/10511937
Publication No. US20060088836A1
GENERAL INFORMATION:
APPLICANT: WOHLGEMUTH, Jay
APPLICANT: Fry, Kirk
APPLICANT: Woodward, Robert
APPLICANT: Ly, Ngoc
APPLICANT: Prentice, James
APPLICANT: Morris, Macdonald
APPLICANT: Rosenberg, Steven
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
AND MONITORING TRANSPLANT REJECTION
FILE REFERENCE: 506612000104
CURRENT APPLICATION NUMBER: US/10/511,937
CURRENT FILING DATE: 2004-10-19
PRIOR APPLICATION NUMBER: PCT/US2003/012946
PRIOR FILING DATE: 2003-04-24
PRIOR APPLICATION NUMBER: US 10/131,831
PRIOR FILING DATE: 2002-04-24
PRIOR APPLICATION NUMBER: US 10/325,899
PRIOR FILING DATE: 2002-12-20
NUMBER OF SEQ ID NOS: 3117
SOFTWARE: PatentIn version 3.2
SEQ ID NO 2476
LENGTH: 153
TYPE: PRT
ORGANISM: Homo sapiens
US-10-511-937-2476

Query Match 66.0%; Score 31; DB 8; Length 153;
Best Local Similarity 75.0%; Pred. No. 37;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
|||||
Db 47 LCTELTVT 54

RESULT 48

US-11-174-398-8
Sequence 8, Application US/11174398
Publication No. US20050244930A1
GENERAL INFORMATION:
APPLICANT: Presnell, Scott R.
APPLICANT: West, James W.
APPLICANT: Novak, Julia E.
TITLE OF INVENTION: ZALPHAL1 LIGAND ANTAGONISTS
FILE REFERENCE: 01-37
CURRENT APPLICATION NUMBER: US/11/174,398
CURRENT FILING DATE: 2005-07-01
PRIOR APPLICATION NUMBER: US/10/282,622
PRIOR FILING DATE: 2002-10-28
PRIOR APPLICATION NUMBER: 60/337,586
PRIOR FILING DATE: 2001-11-05
NUMBER OF SEQ ID NOS: 30
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 8
LENGTH: 153
TYPE: PRT
ORGANISM: Homo sapiens
US-11-174-398-8

Query Match 66.0%; Score 31; DB 11; Length 153;
Best Local Similarity 75.0%; Pred. No. 37;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
| | | | |
| | | | |
Db 47 LCTELTVT 54

RESULT 49
US-11-289-226-13
; Sequence 13, Application US/11289226
; Publication No. US20060084623A1
; GENERAL INFORMATION:
; APPLICANT: Steinman, Lawrence
; APPLICANT: Ruiz, Pedro
; APPLICANT: Garren, Hideki
; TITLE OF INVENTION: DNA Vaccination for Treatment of
; TITLE OF INVENTION: Autoimmune Disease
; FILE REFERENCE: STAN123CTP
; CURRENT APPLICATION NUMBER: US/11/289,226
; CURRENT FILING DATE: 2005-11-28
; PRIOR APPLICATION NUMBER: US/09/947,770
; PRIOR FILING DATE: 2001-09-05
; PRIOR APPLICATION NUMBER: PCT/US00/06233
; PRIOR FILING DATE: 2000-03-10
; PRIOR APPLICATION NUMBER: US 09/267,590
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: amino acid sequence of a Th2 cytokine
US-11-289-226-13

Query Match 66.0%; Score 31; DB 11; Length 153;
Best Local Similarity 75.0%; Pred. No. 37;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
| | | | |
| | | | |
Db 47 LCTELTVT 54

RESULT 50
US-10-530-253-19
; Sequence 19, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaselli, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 19
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 39
US-10-530-253-19

Query Match 66.0%; Score 31; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 38;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 LCTELQTT 9
| | | | |
| | | | |
Db 17 LCTTLDTT 24

Search completed: May 5, 2006, 08:51:43
Job time : 9.4 secs

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OM protein - protein search, using SW model

Run on: May 5, 2006, 06:24:27 ; Search time 26.75 Seconds
(without alignments)
27.816 Million cell updates/sec

Title: US-08-170-344-4
Perfect score: 46
Sequence: 1 LCTELQTTI 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Issued_Patents_AA:*
1: /cgn2_6/pdata/1/1aa/5-COMB.pep:*
2: /cgn2_6/pdata/1/1aa/6-COMB.pep:*
3: /cgn2_6/pdata/1/1aa/H-COMB.pep:*
4: /cgn2_6/pdata/1/1aa/BCITUS-COMB.pep:*
5: /cgn2_6/pdata/1/1aa/RE-COMB.pep:*
6: /cgn2_6/pdata/1/1aa/Backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	46	100.0	20	1	US-08-934-915-159
2	46	100.0	30	1	US-08-363-586-4
3	46	100.0	30	2	US-09-980-523A-4
4	46	100.0	151	2	US-09-701-080C-18
5	46	100.0	158	2	US-09-980-523A-2
6	46	100.0	162	1	US-08-316-239B-3
7	46	100.0	162	1	US-08-316-239B-4
8	46	100.0	172	2	US-08-860-165-14
9	46	100.0	172	2	US-09-359-382-14
10	46	100.0	182	1	US-08-117-083-10
11	46	100.0	243	2	US-09-462-993-1
12	46	100.0	266	2	US-08-860-165-10
13	46	100.0	266	2	US-09-359-382-10
14	46	100.0	266	2	US-09-367-309A-1
15	46	100.0	273	2	US-09-485-885-4
16	46	100.0	292	2	US-09-485-885-10
17	46	100.0	371	2	US-09-485-885-6
18	46	100.0	390	2	US-09-485-885-14
19	46	100.0	390	2	US-08-466-285-2
20	35	76.1	32	1	US-08-164-768-2
21	35	76.1	158	1	US-08-247-904B-10
22	35	76.1	158	2	US-08-767-942A-19
23	35	76.1	271	1	US-08-117-083-14
24	35	76.1	278	2	US-09-485-885-21
25	35	76.1	383	2	US-09-485-885-23
26	34	73.9	637	2	US-09-187-906-7
27	34	73.9	637	2	US-09-489-407-7

28	33	71.7	14	1	US-07-909-122-3	Sequence 3, Appli
29	33	71.7	141	2	US-09-270-767-33527	Sequence 33527, A
30	33	71.7	141	2	US-09-270-767-48744	Sequence 48744, A
31	33	71.7	346	2	US-09-543-681A-6716	Sequence 6716, Ap
32	32	69.6	236	2	US-09-634-137-32	Sequence 32, Appli
33	32	69.6	251	2	US-09-270-767-33015	Sequence 33015, A
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256	28	60.9	416	2	US-09-107-532A-6642	Sequence 6642, Ap	329	28	60.9	1863	2	US-09-462-401A-2	Sequence 2, Appli
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275	28	60.9	765	1	US-08-425-061-19	Sequence 19, Appli	348	27	58.7	21	2	US-09-142-569-17	Sequence 516, App
276	28	60.9	765	1	US-08-825-888-19	Sequence 19, Appli	349	27	58.7	21	2	US-09-495-448A-17	Sequence 437, App
277	28	60.9	765	1	US-08-989-890-19	Sequence 19, Appli	350	27	58.7	36	2	US-09-973-278-437	Sequence 12307, A
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280	28	60.9	900	1	US-08-825-888-20	Sequence 20, Appli	353	27	58.7	60	2	US-09-513-999C-6532	Sequence 14, Appli
281	28	60.9	900	2	US-08-989-890-20	Sequence 21, Appli	354	27	58.7	60	2	US-10-179-784-14	Sequence 40422, A
282	28	60.9	914	1	US-08-425-061-21	Sequence 21, Appli	355	27	58.7	61	2	US-09-270-767-40422	Sequence 55538, A
283	28	60.9	914	1	US-08-825-888-21	Sequence 21, Appli	356	27	58.7	61	2	US-09-248-796A-14272	Sequence 18035, A
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285	28	60.9	940	2	US-09-512-250C-2	Sequence 2, Appli	358	27	58.7	75	2	US-09-540-236-2179	Sequence 40176, A
286	28	60.9	968	2	US-09-949-016-11229	Sequence 11229, A	359	27	58.7	84	2	US-09-270-767-55392	Sequence 13501, A
287	28	60.9	1098	2	US-10-104-047-2475	Sequence 2475, Ap	360	27	58.7	84	2	US-09-902-540-13501	Sequence 48097, A
288	28	60.9	1104	2	US-09-793-998-11	Sequence 11, Appli	361	27	58.7	84	2	US-09-762-724-16	Sequence 41252, A
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291	28	60.9	1202	2	US-08-989-890-22	Sequence 22, Appli	364	27	58.7	90	2	US-09-252-991A-21264	Sequence 44630, A
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294	28	60.9	1363	1	US-08-825-888-23	Sequence 23, Appli	367	27	58.7	107	2	US-09-315-793-42	Sequence 7243, Ap
295	28	60.9	1363	2	US-08-989-890-23	Sequence 6796, Ap	368	27	58.7	125	2	US-09-328-335-1743	Sequence 4, Appli
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297	28	60.9	1598	2	US-09-949-016-6795	Sequence 6795, Ap	370	27	58.7	140	2	US-09-270-767-33783	Sequence 33783, A
298	28	60.9	1634	2	US-09-949-016-6794	Sequence 6794, Ap	371	27	58.7	141	2	US-09-270-767-48000	Sequence 49000, A
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316	28	60.9	1853	1	US-08-825-888-16	Sequence 16, Appli	389	27	58.7				
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409	27	58.7	308	2	US-10-062-548-60	Sequence 60, Appl
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414	27	58.7	322	2	US-09-712-363-212	Sequence 212, App
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417	27	58.7	337	2	US-09-188-930-186	Sequence 186, App
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431	27	58.7	421	2	US-09-948-016-5872	Sequence 5872, Ap
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465	27	58.7	734	2	US-09-949-016-6415	Sequence 6415, Ap
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470	27	58.7	942	2	US-09-171-937C-40	Sequence 40, Appl
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474	27	58.7	1082	2	US-09-762-724-6	Sequence 6, Appli
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481	27	58.7	1650	2	US-09-949-016-11387	Sequence 11387, A
482	27	58.7	1711	2	US-09-771-161A-219	Sequence 219, App
483	27	58.7	1711	2	US-09-771-161A-220	Sequence 220, App
484	27	58.7	2008	2	US-09-091-501B-8	Sequence 8, Appli
485	27	58.7	3433	2	US-09-091-501B-10	Sequence 10, Appl
486	27	58.7	3433	2	US-09-538-092-1136	Sequence 1136, Ap
487	27	58.7	3542	2	US-10-087-013-2	Sequence 2, Appli
488	27	58.7	3696	2	US-09-134-001C-5080	Sequence 5080, Ap
489	27	58.7	10182	2	US-09-134-001C-3159	Sequence 3159, Ap
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493	26	56.5	20	1	US-08-218-025A-40	Sequence 40, Appl
494	26	56.5	24	2	US-09-177-249-197	Sequence 197, App
495	26	56.5	24	2	US-09-812-283-197	Sequence 197, App
496	26	56.5	26	2	US-09-842-164A-28	Sequence 28, Appl
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509	26	56.5	94	2	US-09-270-767-57396	Sequence 57396, A
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523	26	56.5	147	1	US-08-530-165-3	Sequence 3, Appli
524	26	56.5	147	1	US-09-252-991A-26480	Sequence 26480, A
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527	26	56.5	151	2	US-09-934-465-11	Sequence 11, Appl
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535	26	56.5	164	2	US-09-710-279-1266	Sequence 1266, Ap
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537	26	56.5	177	2	US-09-270-767-31838	Sequence 31838, A
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542	26	56.5	194	2	US-09-270-767-47006	Sequence 47006, A	615	26	56.5	359	2	US-09-842-164A-9	Sequence 4, Appl1
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548	26	56.5	213	2	US-09-107-553A-3797	Sequence 3797, Ap	621	26	56.5	384	2	US-09-270-767-52662	Sequence 6, Appl1
549	26	56.5	213	2	US-09-270-767-36590	Sequence 36590, A	622	26	56.5	386	1	US-08-968-751-6	Sequence 6, Appl1
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552	26	56.5	214	1	US-09-094-212-1	Sequence 1, Appl1	625	26	56.5	401	2	US-09-248-796A-16018	Sequence 16018, A
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554	26	56.5	215	1	US-08-462-169B-16	Sequence 16, Appl	627	26	56.5	413	2	US-10-067-443-4	Sequence 4, Appl1
555	26	56.5	215	1	US-08-867-471-6	Sequence 6, Appl1	628	26	56.5	421	2	US-10-067-443-4	Sequence 4, Appl1
556	26	56.5	215	1	US-08-438-438C-6	Sequence 6, Appl1	629	26	56.5	423	2	US-09-243-374-6	Sequence 6, Appl1
557	26	56.5	215	1	US-08-438-438C-16	Sequence 16, Appl	630	26	56.5	442	6	US-08-67-6	Sequence 2, Appl1
558	26	56.5	215	2	US-09-103-079-16	Sequence 16, Appl	631	26	56.5	443	1	US-08-833-963C-2	Sequence 1, Appl1
559	26	56.5	215	2	US-08-705-245-17	Sequence 17, Appl	632	26	56.5	443	2	US-08-980-514-1	Sequence 1, Appl1
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561	26	56.5	215	2	US-09-057-860A-4	Sequence 4, Appl1	634	26	56.5	444	1	US-08-476-008-51	Sequence 51, Appl
562	26	56.5	215	2	US-09-417-721-12	Sequence 12, Appl	635	26	56.5	444	1	US-08-476-008-52	Sequence 52, Appl
563	26	56.5	215	2	US-09-425-021-16	Sequence 16, Appl	636	26	56.5	444	1	US-08-476-008-53	Sequence 53, Appl
564	26	56.5	215	2	US-09-449-249-17	Sequence 17, Appl	637	26	56.5	444	1	US-08-476-008-54	Sequence 54, Appl
565	26	56.5	215	2	US-09-564-829-10	Sequence 10, Appl	638	26	56.5	444	1	US-08-476-008-55	Sequence 55, Appl
566	26	56.5	215	2	US-09-490-714-17	Sequence 17, Appl	639	26	56.5	444	1	US-08-306-063-51	Sequence 51, Appl
567	26	56.5	216	1	US-08-106-507-10	Sequence 10, Appl	640	26	56.5	444	1	US-08-306-063-52	Sequence 52, Appl
568	26	56.5	216	1	US-08-446-922-8	Sequence 8, Appl1	641	26	56.5	444	1	US-08-306-063-53	Sequence 53, Appl
569	26	56.5	216	1	US-08-446-922-8	Sequence 8, Appl1	642	26	56.5	444	1	US-08-306-063-54	Sequence 54, Appl
570	26	56.5	223	6	PCT-US93-10034-8	Sequence 8, Appl1	643	26	56.5	444	1	US-08-306-063-55	Sequence 55, Appl
571	26	56.5	226	2	US-09-489-039A-7406	Sequence 7406, Ap	644	26	56.5	444	1	US-08-306-063-55	Sequence 55, Appl
572	26	56.5	234	2	US-09-248-796A-24043	Sequence 24043, A	645	26	56.5	444	1	US-08-833-485-51	Sequence 51, Appl
573	26	56.5	243	2	US-09-270-767-40663	Sequence 40663, A	646	26	56.5	444	1	US-08-833-485-52	Sequence 52, Appl
574	26	56.5	243	2	US-09-270-767-55879	Sequence 55879, A	647	26	56.5	444	1	US-08-833-485-53	Sequence 53, Appl
575	26	56.5	249	2	US-09-489-039A-8345	Sequence 8345, Ap	648	26	56.5	444	1	US-08-833-485-54	Sequence 54, Appl
576	26	56.5	257	2	US-09-270-767-34358	Sequence 34358, A	649	26	56.5	444	1	US-08-833-485-55	Sequence 55, Appl
577	26	56.5	257	2	US-09-270-767-49575	Sequence 49575, A	650	26	56.5	444	1	US-09-243-374-1	Sequence 1, Appl1
578	26	56.5	262	2	US-09-902-540-12908	Sequence 12908, A	651	26	56.5	444	2	US-09-243-374-2	Sequence 2, Appl1
579	26	56.5	262	2	US-09-710-262B-17	Sequence 17, Appl	652	26	56.5	444	2	US-09-137-440-51	Sequence 51, Appl
580	26	56.5	266	2	US-09-540-236-2940	Sequence 2940, Ap	653	26	56.5	444	2	US-09-137-440-52	Sequence 52, Appl
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582	26	56.5	271	1	US-08-956-012-3	Sequence 3, Appl1	655	26	56.5	444	2	US-09-137-440-54	Sequence 54, Appl
583	26	56.5	271	1	US-10-037-417-26	Sequence 26, Appl	656	26	56.5	444	2	US-09-137-440-55	Sequence 55, Appl
584	26	56.5	274	2	US-09-188-930-336	Sequence 336, App	657	26	56.5	444	2	US-09-137-440-55	Sequence 55, Appl
585	26	56.5	274	2	US-09-312-283C-336	Sequence 336, App	658	26	56.5	444	6	US-09-243-374-4	Sequence 4, Appl1
586	26	56.5	282	2	US-09-540-236-3201	Sequence 3201, Ap	659	26	56.5	446	2	US-09-243-374-4	Sequence 4, Appl1
587	26	56.5	283	2	US-09-134-001C-5534	Sequence 5534, Ap	660	26	56.5	446	6	US-09-243-374-4	Sequence 4, Appl1
588	26	56.5	304	2	US-10-432-803-2	Sequence 2, Appl1	661	26	56.5	446	6	US-09-243-374-4	Sequence 4, Appl1
589	26	56.5	306	2	US-09-252-991A-30319	Sequence 30319, A	662	26	56.5	449	2	US-09-640-419C-24	Sequence 24, Appl
590	26	56.5	306	2	US-10-037-417-89	Sequence 89, Appl	663	26	56.5	449	2	US-09-949-016-7792	Sequence 7792, Ap
591	26	56.5	306	2	US-10-037-417-90	Sequence 90, Appl	664	26	56.5	466	2	US-09-949-016-7792	Sequence 7792, Ap
592	26	56.5	306	4	PCT-US94-09799-1	Sequence 1, Appl1	665	26	56.5	476	2	US-09-489-039A-11933	Sequence 11933, A
593	26	56.5	307	1	US-08-807-861A-56	Sequence 56, Appl	666	26	56.5	482	2	US-10-104-047-3837	Sequence 3837, Ap
594	26	56.5	307	2	US-09-210-681-56	Sequence 56, Appl	667	26	56.5	489	2	US-10-104-047-3922	Sequence 2922, Ap
595	26	56.5	307	2	US-08-946-719A-56	Sequence 56, Appl	668	26	56.5	497	2	US-09-949-016-6348	Sequence 6348, Ap
596	26	56.5	307	2	US-09-747-983-56	Sequence 56, Appl	669	26	56.5	498	2	US-09-270-767-46538	Sequence 46538, A
597	26	56.5	307	2	US-09-743-847-3	Sequence 3, Appl1	670	26	56.5	498	2	US-09-232-468A-18	Sequence 18, Appl
598	26	56.5	307	2	US-10-009-962-7	Sequence 7, Appl1	671	26	56.5	498	2	US-08-686-968C-231	Sequence 231, App
599	26	56.5	307	2	US-10-037-417-87	Sequence 87, Appl	672	26	56.5	501	1	US-08-448-603A-2	Sequence 2, Appl1
600	26	56.5	307	2	US-10-037-417-88	Sequence 88, Appl	673	26	56.5	501	2	US-09-134-075-2	Sequence 2, Appl1
601	26	56.5	309	2	US-10-671-628-8	Sequence 8, Appl	674	26	56.5	501	2	US-09-492-739-2	Sequence 2, Appl1
602	26	56.5	309	2	US-09-449-016-10401	Sequence 7, Appl1	675	26	56.5	501	2	US-09-966-931A-2	Sequence 2, Appl1
603	26	56.5	311	2	US-09-543-681A-5602	Sequence 10401, A	676	26	56.5	507	2	US-09-949-016-10371	Sequence 10371, A
604	26	56.5	312	2	US-09-242-948-4	Sequence 5602, Ap	677	26	56.5	511	1	US-08-448-603A-1	Sequence 1, Appl1
605	26	56.5	315	2	US-08-793-035-9	Sequence 4, Appl1	678	26	56.5	511	2	US-09-134-075-5	Sequence 5, Appl1
606	26	56.5	315	2	US-08-793-035-10	Sequence 9, Appl1	679	26	56.5	511	2	US-08-889-841B-41	Sequence 41, Appl
607	26	56.5	317	1	US-08-767-096-2	Sequence 10, Appl	680	26	56.5	511	2	US-09-492-739-1	Sequence 1, Appl1
608	26	56.5	317	2	US-09-480-203-2	Sequence 2, Appl1	681	26	56.5	511	2	US-09-419-362-41	Sequence 41, Appl
609	26	56.5	327	2	US-09-543-681A-4517	Sequence 4517, Ap	682	26	56.5	513	2	US-09-966-931A-1	Sequence 1, Appl1
610	26	56.5	327	2	US-10-104-047-2508	Sequence 2508, Ap	683	26	56.5	513	2	US-08-472-240A-14	Sequence 14, Appl
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686	26	56.5	516	2	US-09-685-403-27	Sequence 27, Appl	759	26	56.5	1125	2	US-09-900-920-60	Sequence 60, Appl
687	26	56.5	516	6	5188642-4	Patent No. 5188642	760	26	56.5	1125	2	US-09-487-558B-106	Sequence 106, App
688	26	56.5	520	2	US-09-685-403-2	Sequence 2, Appl1	761	26	56.5	1252	2	US-09-902-540-13967	Sequence 13967, A
689	26	56.5	530	2	US-08-979-608A-8	Sequence 8, Appl1	762	26	56.5	1388	2	US-08-685-576-1	Sequence 1, Appl1
690	26	56.5	530	2	US-09-517-848-8	Sequence 8, Appl1	763	26	56.5	1648	1	US-09-515-806-4	Sequence 4, Appl1
691	26	56.5	530	2	US-09-616-289-8	Sequence 8, Appl1	764	26	56.5	1778	2	US-09-252-991A-18159	Sequence 18159, A
692	26	56.5	530	2	US-09-976-740-8	Sequence 8, Appl1	765	26	56.5	1979	2	US-09-949-016-6468	Sequence 6468, Ap
693	26	56.5	541	2	US-09-716-129-54	Sequence 54, Appl	766	26	56.5	2047	2	US-09-949-016-7404	Sequence 7404, Ap
694	26	56.5	543	2	US-09-270-767-59648	Sequence 59648, A	767	26	56.5	2221	2	US-10-144-198-30	Sequence 30, Appl
695	26	56.5	543	2	US-09-351-150A-15	Sequence 15, Appl	768	26	56.5	2930	2	US-09-417-822-2	Sequence 2, Appl1
696	26	56.5	546	2	US-09-616-289-44	Sequence 44, Appl	769	26	56.5	2930	2	US-09-957-837A-2	Sequence 2, Appl1
697	26	56.5	546	2	US-09-976-740-44	Sequence 44, Appl	770	26	56.5	3224	1	US-08-705-660-14	Sequence 34, Appl
698	26	56.5	550	1	US-08-417-210A-140	Sequence 140, App	771	26	56.5	3224	2	US-08-989-046-14	Sequence 34, Appl
699	26	56.5	550	1	US-09-136-159A-140	Sequence 140, App	772	26	56.5	3224	2	US-09-538-092-1161	Sequence 1161, Ap
700	26	56.5	551	1	US-08-417-210A-137	Sequence 137, App	773	26	56.5	3224	2	US-09-315-335A-34	Sequence 34, Appl
701	26	56.5	551	1	US-08-417-210A-137	Sequence 137, App	774	25.5	55.4	56	2	US-09-621-976-6978	Sequence 6978, Ap
702	26	56.5	551	2	US-09-136-159A-137	Sequence 137, App	775	25.5	55.4	3170	2	US-09-036-987A-4	Sequence 4, Appl1
703	26	56.5	551	2	US-09-136-159A-143	Sequence 143, App	776	25.5	55.4	3170	2	US-09-370-700-4	Sequence 4, Appl1
704	26	56.5	555	2	US-09-270-767-61186	Sequence 61186, A	777	25.5	55.4	3170	2	US-09-603-207-4	Sequence 4, Appl1
705	26	56.5	557	2	US-08-979-608A-5	Sequence 5, Appl1	778	25	54.3	9	1	US-08-629-291A-27	Sequence 27, Appl
706	26	56.5	557	2	US-09-517-849-5	Sequence 5, Appl1	779	25	54.3	9	1	US-08-658-335B-27	Sequence 27, Appl
707	26	56.5	557	2	US-09-616-289-5	Sequence 5, Appl1	780	25	54.3	9	2	US-08-159-339A-329	Sequence 329, App
708	26	56.5	557	2	US-09-976-740-5	Sequence 5, Appl1	781	25	54.3	9	2	US-09-406-640-27	Sequence 27, Appl
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710	26	56.5	615	2	US-09-107-532A-6507	Sequence 6507, Ap	783	25	54.3	10	2	US-08-159-339A-559	Sequence 559, App
711	26	56.5	619	2	US-09-248-796A-17705	Sequence 17705, A	784	25	54.3	11	2	US-09-601-729-176	Sequence 176, App
712	26	56.5	622	2	US-10-363-937-11	Sequence 11, Appl	785	25	54.3	17	2	US-09-525-269A-14	Sequence 14, Appl
713	26	56.5	629	2	US-09-270-767-44226	Sequence 44226, A	786	25	54.3	19	6	5336758-1	Patent No. 5336758
714	26	56.5	655	2	US-09-949-016-9577	Sequence 9577, Ap	787	25	54.3	24	2	US-09-461-697-177	Sequence 12, Appl
715	26	56.5	664	2	US-10-181-612-2	Sequence 2, Appl1	788	25	54.3	25	1	US-08-620-151-23	Sequence 23, App
716	26	56.5	688	2	US-09-949-016-10779	Sequence 10779, A	789	25	54.3	31	1	US-08-438-753B-7	Sequence 7, Appl
717	26	56.5	692	2	US-09-252-991A-19668	Sequence 19668, A	790	25	54.3	31	1	US-08-438-753B-7	Sequence 7, Appl
718	26	56.5	730	2	US-09-248-796A-19567	Sequence 19567, A	791	25	54.3	31	1	US-08-631-328-7	Sequence 7, Appl
719	26	56.5	768	2	US-09-302-812-8	Sequence 8, Appl1	792	25	54.3	31	1	US-08-455-524B-7	Sequence 7, Appl1
720	26	56.5	768	2	US-09-511-477-8	Sequence 8, Appl1	793	25	54.3	31	1	US-08-455-021B-7	Sequence 7, Appl1
721	26	56.5	768	2	US-09-511-507-8	Sequence 8, Appl1	794	25	54.3	31	1	US-09-045-467-7	Sequence 7, Appl1
722	26	56.5	782	2	US-09-585-858-29	Sequence 29, Appl	795	25	54.3	32	1	US-07-952-735A-3	Sequence 3, Appl1
723	26	56.5	782	2	US-10-270-878-29	Sequence 29, Appl	796	25	54.3	34	1	US-07-952-735A-1	Sequence 1, Appl1
724	26	56.5	834	1	US-08-491-357-2	Sequence 2, Appl1	797	25	54.3	34	1	US-07-952-735A-2	Sequence 2, Appl1
725	26	56.5	834	1	US-08-968-633-2	Sequence 2, Appl1	798	25	54.3	37	1	US-07-946-054-1	Sequence 1, Appl1
726	26	56.5	834	2	US-09-196-466-2	Sequence 2, Appl1	799	25	54.3	37	1	US-08-262-037-3	Sequence 3, Appl1
727	26	56.5	834	2	US-09-669-459A-2	Sequence 2, Appl1	800	25	54.3	37	4	PCT-US93-08638-1	Sequence 1, Appl1
728	26	56.5	834	4	PCT-US96-10823-2	Sequence 2, Appl1	801	25	54.3	48	2	US-09-902-540-12332	Sequence 12332, A
729	26	56.5	856	1	US-08-022-835-2	Sequence 2, Appl1	802	25	54.3	49	1	US-08-417-210A-114	Sequence 114, App
730	26	56.5	856	1	US-08-388-809-2	Sequence 2, Appl1	803	25	54.3	49	2	US-09-136-159A-114	Sequence 114, App
731	26	56.5	856	1	US-08-647-714-2	Sequence 2, Appl1	804	25	54.3	58	2	US-09-627-775-10	Sequence 10, Appl
732	26	56.5	856	1	US-07-956-483-11	Sequence 11, Appl	805	25	54.3	59	1	US-08-470-179-4	Sequence 4, Appl1
733	26	56.5	857	1	US-08-022-835-4	Sequence 4, Appl1	806	25	54.3	62	2	US-09-902-540-16179	Sequence 16179, A
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735	26	56.5	857	1	US-08-647-714-4	Sequence 4, Appl1	808	25	54.3	74	2	US-09-525-269A-1	Sequence 1, Appl1
736	26	56.5	877	2	US-08-612-353-2	Sequence 2, Appl1	809	25	54.3	74	2	US-09-525-269A-2	Sequence 2, Appl1
737	26	56.5	887	2	US-09-472-240A-2	Sequence 2, Appl1	810	25	54.3	75	1	US-08-428-415-22	Sequence 22, Appl
738	26	56.5	891	2	US-10-226-629A-16	Sequence 16, Appl	811	25	54.3	75	1	US-08-379-685-22	Sequence 22, Appl
739	26	56.5	892	2	US-10-226-629A-15	Sequence 15, Appl	812	25	54.3	75	1	US-08-854-029-22	Sequence 22, Appl
740	26	56.5	897	2	US-09-543-681A-4249	Sequence 4249, Ap	813	25	54.3	75	2	US-08-428-762-22	Sequence 22, Appl
741	26	56.5	903	2	US-09-489-039A-13692	Sequence 13692, A	814	25	54.3	75	2	US-09-248-796A-26588	Sequence 26588, A
742	26	56.5	903	1	US-08-673-789-7	Sequence 7, Appl1	815	25	54.3	76	2	US-09-083-521-5	Sequence 5, Appl1
743	26	56.5	970	1	US-08-162-809-10	Sequence 10, Appl	816	25	54.3	77	2	US-09-248-796A-23996	Sequence 23996, A
744	26	56.5	980	2	US-09-627-376-7	Sequence 7, Appl1	817	25	54.3	99	2	US-09-487-558B-26392	Sequence 26392, A
745	26	56.5	980	2	US-09-442-100-8	Sequence 8, Appl1	818	25	54.3	88	2	US-09-270-767-40342	Sequence 40342, A
746	26	56.5	980	2	US-08-939-106-8	Sequence 8, Appl1	819	25	54.3	89	2	US-09-270-767-55558	Sequence 55558, A
747	26	56.5	980	2	US-09-442-102-8	Sequence 8, Appl1	820	25	54.3	90	2	US-09-107-532A-6712	Sequence 6712, Ap
748	26	56.5	983	2	US-09-538-092-1320	Sequence 1320, Ap	821	25	54.3	99	2	US-09-270-767-62266	Sequence 62266, A
749	26	56.5	988	2	US-08-162-809-14	Sequence 14, Appl	822	25	54.3	102	1	US-08-313-185-64	Sequence 64, Appl
750	26	56.5	990	2	US-09-627-376-7	Sequence 7, Appl1	823	25	54.3	102	1	US-08-808-982-8	Sequence 8, Appl1
751	26	56.5	990	2	US-10-047-676B-7	Sequence 7, Appl1	824	25	54.3	102	1	US-09-306-902A-8	Sequence 8, Appl1
752	26	56.5	1002	2	US-09-252-991A-24655	Sequence 24655, A	825	25	54.3	102	2	US-09-270-767-49308	Sequence 49308, A
753	26	56.5	1003	2	US-09-319-558B-6	Sequence 6, Appl1	826	25	54.3	104	2	US-09-487-558B-242	Sequence 242, App
754	26	56.5	1014	2	US-09-487-558B-6	Sequence 6, Appl1	827	25	54.3	105	2	US-09-543-681A-5338	Sequence 5338, Ap
755	26	56.5	1068	2	US-09-136-652-2	Sequence 2, Appl1	828	25	54.3	107	2	US-09-134-001C-2887	Sequence 2887, Ap
756	26	56.5	1079	2	US-09-538-092-812	Sequence 812, App	829	25	54.3				
757	26	56.5	1083	2			830	25	54.3				

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832	25	54.3	108	2	US-09-270-767-50695	Sequence 50695, A	905	25	54.3	172	1	US-08-443-883A-2	Sequence 2, Appl1
833	25	54.3	110	2	US-09-270-767-58542	Sequence 58542, A	906	25	54.3	172	1	US-08-631-328-2	Sequence 2, Appl1
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836	25	54.3	120	2	US-09-488-039A-12045	Sequence 12045, A	909	25	54.3	172	2	US-09-045-467-2	Sequence 2, Appl1
837	25	54.3	123	2	US-09-252-991A-21343	Sequence 21343, A	910	25	54.3	172	2	US-08-954-395A-18	Sequence 18, Appl1
838	25	54.3	123	2	US-09-732-210-1144	Sequence 1144, Ap	911	25	54.3	172	2	US-08-616-904-2	Sequence 2, Appl1
839	25	54.3	123	2	US-09-732-210-1145	Sequence 1145, Ap	912	25	54.3	172	2	US-09-599-413-2	Sequence 2, Appl1
840	25	54.3	123	2	US-09-732-210-1152	Sequence 1152, Ap	913	25	54.3	172	2	US-09-599-413-4	Sequence 5, Appl1
841	25	54.3	123	2	US-09-732-210-1158	Sequence 1158, Ap	914	25	54.3	172	2	US-09-599-413-6	Sequence 6, Appl1
842	25	54.3	123	2	US-09-732-210-1322	Sequence 1322, Ap	915	25	54.3	172	2	US-09-599-413-7	Sequence 7, Appl1
843	25	54.3	123	2	US-09-732-210-1324	Sequence 1324, Ap	916	25	54.3	172	2	US-09-599-413-8	Sequence 8, Appl1
844	25	54.3	123	2	US-09-270-767-32548	Sequence 32548, A	917	25	54.3	172	2	US-09-599-413-9	Sequence 9, Appl1
845	25	54.3	123	2	US-09-270-767-32548	Sequence 47765, A	918	25	54.3	172	2	US-09-599-413-10	Sequence 10, Appl1
846	25	54.3	123	2	US-09-855-266A-13	Sequence 13, Appl1	919	25	54.3	172	2	US-09-599-413-18	Sequence 18, Appl1
847	25	54.3	123	2	US-10-142-835-14	Sequence 14, Appl1	920	25	54.3	172	2	US-09-599-413-19	Sequence 19, Appl1
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851	25	54.3	124	2	US-09-732-210-1323	Sequence 1332, Ap	924	25	54.3	175	2	US-09-270-767-48770	Sequence 5206, Ap
852	25	54.3	124	2	US-09-732-210-1332	Sequence 1332, Ap	925	25	54.3	175	2	US-09-134-000C-5206	Sequence 4398, Ap
853	25	54.3	124	2	US-09-732-210-1335	Sequence 1335, Ap	926	25	54.3	176	2	US-09-107-433-4398	Sequence 10368, A
854	25	54.3	124	2	US-09-732-210-1338	Sequence 1339, Ap	927	25	54.3	180	2	US-09-902-540-10368	Sequence 22, Appl1
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863	25	54.3	127	2	US-09-887-052-8	Sequence 8, Appl1	936	25	54.3	184	2	US-09-408-020-32	Sequence 9063, Ap
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875	25	54.3	140	2	US-09-617-302-5	Sequence 46, Appl1	948	25	54.3	195	2	US-09-908-594-11	Sequence 11, Appl1
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877	25	54.3	140	2	US-09-886-319A-48	Sequence 18, Appl1	950	25	54.3	196	2	US-09-908-594-12	Sequence 12, Appl1
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880	25	54.3	141	2	US-08-906-616-135	Sequence 135, App	953	25	54.3	197	2	US-09-902-540-12806	Sequence 4, Appl1
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886	25	54.3	144	2	US-09-770-767-41629	Sequence 41629, A	959	25	54.3	207	2	US-09-252-991A-1993	Sequence 683, Ap
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994 25 54.3 245 2 US-09-489-039A-9463
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ALIGNMENTS

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Sequence 10609, A
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Sequence 29, Appl
Sequence 33, Appl
Sequence 9463, Ap
Sequence 429, App
Sequence 23, Appl
Sequence 23, Appl
Sequence 31, Appl
Sequence 34, Appl
Sequence 6621, Ap

RESULT 1
US-08-934-915-159
; Sequence 159, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
; TITLE OF INVENTION: DIAGNOSTIC PURPOSES
; NUMBER OF SEQUENCES: 193
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft Word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. FOUTCH
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946,6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:
; INFORMATION FOR SEQ ID NO: 159:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid

TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-934-915-159

Query Match 100.0%; Score 46; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 0.048;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 6 LCTELQTTI 14

RESULT 2
US-08-363-586-4
; Sequence 4, Application US/08363586
; Patent No. 5629161
; GENERAL INFORMATION:
; APPLICANT: Mueller, Martin
; APPLICANT: Giesmann, Lutz
; TITLE OF INVENTION: Use of HPV-16 E6 and E7-gene Derived
; TITLE OF INVENTION: Peptides for the Diagnostic Purpose
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; ADDRESS: Dunner
; STREET: 1300 I Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/363,586
; FILING DATE: 23-DEC-1994
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/909,296
; FILING DATE: 09-JUL-1992
; APPLICATION NUMBER: EP 91111720.8
; FILING DATE: 13-JUL-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Wadler, Linda A.
; REGISTRATION NUMBER: 33,218
; REFERENCE/DOCKET NUMBER: 02481-1195-00000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-408-4000
; TELEFAX: 202-408-4400
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 30 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-363-586-4

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Best Local Similarity 100.0%; Pred. No. 0.072;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 3
US-09-980-523A-4
; Sequence 4, Application US/09980523A
; Patent No. 6783763


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; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-4

Query Match          100.0%; Score 46; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.072;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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Db 8 LCTBLQTTI 16

RESULT 4
US-09-701-080C-18
; Sequence 18, Application US/09701080C
; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; CURRENT FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match          100.0%; Score 46; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.36;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTBLQTTI 9
Db 15 LCTBLQTTI 23

RESULT 5
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
```

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; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match          100.0%; Score 46; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.38;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTBLQTTI 9
Db 22 LCTBLQTTI 30

RESULT 6
US-08-316-239B-3
; Sequence 3, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmentier, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESS: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
US-08-316-239B-3
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Query Match 100.0%; Score 46; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.39; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
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Db 22 LCTELQTTI 30

RESULT 7

US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNNE-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
; US-08-316-239B-4

Query Match 100.0%; Score 46; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.39; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
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Db 22 LCTELQTTI 30

RESULT 8

US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian

; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/1130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE INFORMATION: Description of Artificial Sequence: Gene Fusion
; US-08-860-165-14

Query Match 100.0%; Score 46; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.41; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
| | | | | | | | | |
Db 91 LCTELQTTI 99

RESULT 9

US-09-359-382-14
; Sequence 14, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
; US-09-359-382-14

Query Match 100.0%; Score 46; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.41; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
| | | | | | | | | |
Db 91 LCTELQTTI 99

RESULT 10

US-08-117-083-10
; Sequence 10, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Boursnell, Michael E.
; APPLICANT: Inglis, Stephen C.
; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; TITLE OF INVENTION: Papilloma Virus Proteins

```

;
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Walter H. Dreger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dreger, Walter H.
; REGISTRATION NUMBER: 24,190
; REFERENCE/DOCKET NUMBER: A-58783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 182 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..182
; OTHER INFORMATION: /note= "Xaa refers to stop codon in
; OTHER INFORMATION: the open reading frame."
;
; US-08-117-083-10
;
Query Match 100.0%; Score 46; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.43;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
;
Qy 1 LCTELQTTI 9
Db 23 LCTELQTTI 31
;
RESULT 11
; US-09-462-993-1
; Sequence 1, Application US/09462993
; Patent No. 6884786
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BALLOU, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/09/462,993
; CURRENT FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Derived from

```

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;
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone E6*TMF.
;
; US-09-462-993-1
;
Query Match 100.0%; Score 46; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.58;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
;
Qy 1 LCTELQTTI 9
Db 50 LCTELQTTI 58
;
RESULT 12
; US-08-860-165-10
; Sequence 10, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
;
; US-08-860-165-10
;
Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.63;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
;
Qy 1 LCTELQTTI 9
Db 22 LCTELQTTI 30
;
RESULT 13
; US-09-359-382-10
; Sequence 10, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PN0157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT

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ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.63;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 22 LCTELQTTI 30

RESULT 14
US-09-367-309A-1
Sequence 1, Application US/09367309A

Patent No. 6428807
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
PRIOR FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.63;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 22 LCTELQTTI 30

RESULT 15
US-09-485-885-4
Sequence 4, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 46; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.65;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 128 LCTELQTTI 136

RESULT 16
US-09-485-885-10
Sequence 10, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 46; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.7;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 147 LCTELQTTI 155

RESULT 17
US-09-485-885-6
Sequence 6, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 46; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.88;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Db 128 LCTELQTTI 136

RESULT 18

US-09-485-885-14
Sequence 14, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Benchehly, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-07-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 46; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.93;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9

Db 147 LCTELQTTI 155

RESULT 19

US-08-466-285-2
Sequence 2, Application US/08466285
Patent No. 5753233
GENERAL INFORMATION:
APPLICANT: Bleul, Conrad
APPLICANT: Giesmann, Lutz
APPLICANT: Muller, Martin
TITLE OF INVENTION: Seroreactive Epitopes On Proteins Of
TITLE OF INVENTION: Human Papillomavirus (HPV) 18
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESS: 1300 I Street, N.W., Suite 700
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,285
FILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/947,992
FILING DATE: 21-SEP-1992

CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/696,953
FILING DATE: 08-MAY-1991
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: P 40 15 044.5
FILING DATE: 10-MAY-1990
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Manpeizer, David A.
REGISTRATION NUMBER: 37,540
REFERENCE/DOCKET NUMBER: 05552.1075-03000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-466-285-2

Query Match 76.1%; Score 35; DB 1; Length 32;
Best Local Similarity 66.7%; Pred. No. 7.9;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9

Db 12 LCTELQTTI 20

RESULT 20

US-08-164-768-2
Sequence 2, Application US/08164768
Patent No. 6322794
GENERAL INFORMATION:
APPLICANT: BLEUL, Conrad
APPLICANT: GISSMANN, Lutz
APPLICANT: MULLER, Martin
TITLE OF INVENTION: SEROREACTIVE EPITOPES ON PROTEINS OF
TITLE OF INVENTION: HUMAN PAPILLOMA VIRUS (HPV) 18
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: FINNEGAN, HENDERSON, FARABOW, GARRETT &
ADDRESS: DUNNER, L.L.P.
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Forman, David S.
REGISTRATION NUMBER: 33,694
REFERENCE/DOCKET NUMBER: 05552.1075-02000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid

STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-164-768-2

Query Match 76.1%; Score 35; DB 2; Length 32;
Best Local Similarity 66.7%; Pred. No. 7.9;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 12 LCTELNTSL 20

RESULT 21
US-08-247-904B-10
Sequence 10, Application US/08247904B
Patent No. 5981699
GENERAL INFORMATION:
APPLICANT: Rolfe, Mark
APPLICANT: Eckstein, Jens W.
TITLE OF INVENTION: Human ubiquitin Conjugating Enzyme
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Foley, Hoag & Eliot
STREET: One Post Office Square
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII(text)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/247,904B
FILING DATE: 23-MAY-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: MIV-029.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 832-1000
TELEFAX: (617) 832-7000
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-247-904B-10

Query Match 76.1%; Score 35; DB 1; Length 158;
Best Local Similarity 66.7%; Pred. No. 39;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 17 LCTELNTSL 25

RESULT 22
US-08-767-942A-19
Sequence 19, Application US/08767942A
Patent No. 6068982
GENERAL INFORMATION:
APPLICANT: Rolfe, Mark
APPLICANT: Chiu, M. Isabel
APPLICANT: Berlin, Vivian
APPLICANT: Damagnez, Veronique

APPLICANT: Draetta, Giulio
APPLICANT: Guillaume, Cottarel
TITLE OF INVENTION: UBIQUITIN CONJUGATING ENZYMES
NUMBER OF SEQUENCES: 45
CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY, HOAG & ELIOT LLP
STREET: One Post Office Square
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02109-2170
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/767,942A
FILING DATE: 17-DEC-1996
ATTORNEY/AGENT INFORMATION:
NAME: Vincent, Matthew P.
REGISTRATION NUMBER: 36,709
REFERENCE/DOCKET NUMBER: MIV-029.04
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-832-1000
TELEFAX: 617-832-7000
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 158 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-767-942A-19

Query Match 76.1%; Score 35; DB 2; Length 158;
Best Local Similarity 66.7%; Pred. No. 39;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 17 LCTELNTSL 25

RESULT 23
US-08-117-083-14
Sequence 14, Application US/08117083
Patent No. 5719054
GENERAL INFORMATION:
APPLICANT: Boursnell, Michael E.
APPLICANT: Inglis, Stephen C.
APPLICANT: Munro, Alan J.
TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSEE: Walter H. Dreger
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,083
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190

```

; REFERENCE/DOCKET NUMBER: A-58783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 272299
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 271 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..271
; OTHER INFORMATION: /note="Xaa refers to stop codon in
; OTHER INFORMATION: the open reading frame."
US-08-117-083-14

Query Match 76.1%; Score 35; DB 1; Length 271;
Best Local Similarity 66.7%; Pred. No. 67;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 18 LCTELNTSL 26

RESULT 24
US-09-485-885-21
; Sequence 21, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 21
; LENGTH: 278
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-21

Query Match 76.1%; Score 35; DB 2; Length 278;
Best Local Similarity 66.7%; Pred. No. 68;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 128 LCTELNTSL 136

RESULT 25
US-09-485-885-23
; Sequence 23, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
```

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; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 23
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-23

Query Match 76.1%; Score 35; DB 2; Length 383;
Best Local Similarity 66.7%; Pred. No. 94;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 128 LCTELNTSL 136

RESULT 26
US-09-187-906-7
; Sequence 7, Application US/09187906
; Patent No. 6677135
; GENERAL INFORMATION:
; APPLICANT: BIOGEN, INC.
; TITLE OF INVENTION: Ret Ligand (RetL) for Stimulating Neural
; TITLE OF INVENTION: and Renal Growth
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESSES:
; ADDRESS: Biogen, Inc.
; STREET: 14 Cambridge Center
; CITY: Cambridge
; STATE: MA
; COUNTRY: USA
; ZIP: 02142
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/187,906
; FILING DATE:
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/US97/07726
; FILING DATE: 07-MAY-97
; APPLICATION NUMBER: US 60/017,427
; FILING DATE: 08-MAY-96
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/019,300
; FILING DATE: 07-JUN-96
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/021,859
; FILING DATE: 16-JUL-96
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 60/043,533
; FILING DATE: 10-APR-97
; ATTORNEY/AGENT INFORMATION:
; NAME: Kaplan, Warren A.
; REGISTRATION NUMBER: 34,199
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-679-2400
; TELEFAX: 617-679-2838
; INFORMATION FOR SEQ ID NO: 7:
; SEQUENCE CHARACTERISTICS:
```

LENGTH: 637 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-187-906-7

Query Match 73.9%; Score 34; DB 2; Length 637;
Best Local Similarity 75.0%; Pred. No. 2.4e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 2 CTELOTTI 9
Db 479 CTELOTTV 486

RESULT 27
US-09-489-407-7
Sequence 7, Application US/09489407
Patent No. 6861509
GENERAL INFORMATION:
APPLICANT: BIOGEN, INC.
TITLE OF INVENTION: Ret Ligand (RetL) for Stimulating Neural
TITLE OF INVENTION: and Renal Growth
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSEE: Biogen, Inc.
STREET: 14 Cambridge Center
CITY: Cambridge
STATE: MA
COUNTRY: USA
ZIP: 02142
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/489,407
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US97/07726
FILING DATE: 07-MAY-97
APPLICATION NUMBER: US 60/017,427
FILING DATE: 08-MAY-96
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/019,300
FILING DATE: 07-JUN-96
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/021,859
FILING DATE: 16-JUL-96
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/043,533
FILING DATE: 10-APR-97
ATTORNEY/AGENT INFORMATION:
NAME: Kaplan, Warren A.
REGISTRATION NUMBER: 34,199
REFERENCE/DOCKET NUMBER: A008 PCT CIP
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-679-2400
TELEFAX: 617-679-2838
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
LENGTH: 637 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-09-489-407-7

Query Match 73.9%; Score 34; DB 2; Length 637;
Best Local Similarity 75.0%; Pred. No. 2.4e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 2 CTELOTTI 9
Db 479 CTELOTTV 486

RESULT 28
US-07-909-122-3
Sequence 3, Application US/07909122
Patent No. 5415995
GENERAL INFORMATION:
APPLICANT: SCHOOLNIK, GARY K.
APPLICANT: PALESKY, JOEL M.
TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
TITLE OF INVENTION: VIRUS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/909,122
FILING DATE: 19920706
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: BENZ, WILLIAM H.
REGISTRATION NUMBER: 25,952
REFERENCE/DOCKET NUMBER: 28600-20105.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
US-07-909-122-3

Query Match 71.7%; Score 33; DB 1; Length 14;
Best Local Similarity 100.0%; Pred. No. 8;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 3 TELQTTI 9
Db 1 TELQTTI 7

RESULT 29
US-09-270-767-33527
Sequence 33527, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 33527
LENGTH: 141
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-33527


```
Query Match          71.7%; Score 33; DB 2; Length 141;
Best Local Similarity 66.7%; Pred. No. 81;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY      1 LCTELOTTI 9
        |||||
        :
Db      46 LCTELOLKV 54

RESULT 30
US-09-270-767-48744
; Sequence 48744, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 48744
; LENGTH: 141
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-48744

Query Match          71.7%; Score 33; DB 2; Length 141;
Best Local Similarity 66.7%; Pred. No. 81;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY      1 LCTELOTTI 9
        |||||
        :
Db      46 LCTELOLKV 54

RESULT 31
US-09-543-681A-6716
; Sequence 6716, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; CURRENT FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO 6716
; LENGTH: 346
; TYPE: PRT
; ORGANISM: Proteus mirabilis
US-09-543-681A-6716

Query Match          71.7%; Score 33; DB 2; Length 346;
Best Local Similarity 55.6%; Pred. No. 2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 LCTELOTTI 9
        |||||
        :
Db      318 MCTOLOPVL 326

RESULT 32
US-09-634-137-32
; Sequence 32, Application US/09634137
; Patent No. 6632665
; GENERAL INFORMATION:
; APPLICANT: Perrino, Fred W
; TITLE OF INVENTION: Mammalian Genes Encoding 3'-5' Exonuclelease
```

```
; FILE REFERENCE: wak200/48001/4-018
; CURRENT APPLICATION NUMBER: US/09/634,137
; CURRENT FILING DATE: 2000-08-08
; PRIOR APPLICATION NUMBER: US 60/148,018
; PRIOR FILING DATE: 1999-08-09
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 32
; LENGTH: 236
; TYPE: PRT
; ORGANISM: Mus musculus
US-09-634-137-32

Query Match          69.6%; Score 32; DB 2; Length 236;
Best Local Similarity 100.0%; Pred. No. 2.1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 LCTELO 6
        |||||
        :
Db      127 LCTELO 132

RESULT 33
US-09-270-767-33015
; Sequence 33015, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 33015
; LENGTH: 251
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-33015

Query Match          69.6%; Score 32; DB 2; Length 251;
Best Local Similarity 66.7%; Pred. No. 2.2e+02;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY      1 LCTELOTTI 9
        |||||
        :
Db      239 LCTTLXTLL 247

RESULT 34
US-08-271-539-3
; Sequence 3, Application US/08271539
; Patent No. 6358509
; GENERAL INFORMATION:
; APPLICANT: Ramanathan, Lata
; APPLICANT: Seelitz, Gail F.
; APPLICANT: Trotta, Paul P.
; TITLE OF INVENTION: Antibody Antagonists of Human Interleukin-4
; FILE REFERENCE: JB0059KQ US
; CURRENT APPLICATION NUMBER: US/08/271,539
; CURRENT FILING DATE: 1994-07-07
; PRIOR APPLICATION NUMBER: US 07/453,570
; PRIOR FILING DATE: 1989-12-20
; PRIOR APPLICATION NUMBER: PCT/US90/07289
; PRIOR FILING DATE: 1990-12-18
; PRIOR APPLICATION NUMBER: US 07/859,689
; PRIOR FILING DATE: 1992-06-11
; NUMBER OF SEQ ID NOS: 46
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 3
; LENGTH: 14
```

;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: Description of Artificial Sequence: synthetic
;; OTHER INFORMATION: polypeptide
US-08-271-539-3

Query Match 67.4%; Score 31; DB 2; Length 14;
Best Local Similarity 75.0%; Pred. No. 19;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 LCTELQTT 8
Db 6 LCTELTYT 13

RESULT 35
US-08-234-812-3
; Sequence 3, Application US/08234812
; Patent No. 5557535
; GENERAL INFORMATION:
; APPLICANT: Srinivasan, Subhashini
; APPLICANT: Sudarsanam, Padmanaban
; TITLE OF INVENTION: METHOD AND SYSTEM FOR PROTEIN MODELING
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Seed and Berry
; STREET: 6300 Columbia Center, 701 Fifth Avenue
; CITY: Seattle
; STATE: Washington
; COUNTRY: US
; ZIP: 98104-7092
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/234,812
; FILING DATE: 28-APR-1994
; CLASSIFICATION: 395
; ATTORNEY/AGENT INFORMATION:
; NAME: Pirlo, Maurice J.
; REGISTRATION NUMBER: 33,273
; REFERENCE/DOCKET NUMBER: 480052.408C1
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 622-4900
; TELEFAX: (206) 682-6031
; TELEX: 3723836 SEEDANBERRY
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 108 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-08-234-812-3

Query Match 67.4%; Score 31; DB 1; Length 108;
Best Local Similarity 75.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 LCTELQTT 8
Db 17 LCTELTYT 24

RESULT 36
US-08-663-809-3
; Sequence 3, Application US/08663809
; Patent No. 5884230
; GENERAL INFORMATION:
; APPLICANT: Srinivasan, Subhashini
; APPLICANT: Sudarsanam, Padmanaban

;; TITLE OF INVENTION: METHOD AND SYSTEM FOR PROTEIN MODELING
;; NUMBER OF SEQUENCES: 3
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Seed and Berry
;; STREET: 6300 Columbia Center, 701 Fifth Avenue
;; CITY: Seattle
;; STATE: Washington
;; COUNTRY: US
;; ZIP: 98104-7092
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Floppy disk
;; COMPUTER: IBM PC compatible
;; OPERATING SYSTEM: PC-DOS/MS-DOS
;; SOFTWARE: Patentin Release #1.0, Version #1.25
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/663,809
;; FILING DATE: 14-JUN-1996
;; CLASSIFICATION: 364
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Pirlo, Maurice J.
;; REGISTRATION NUMBER: 33,273
;; REFERENCE/DOCKET NUMBER: 480052.408C2
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (206) 622-4900
;; TELEFAX: (206) 682-6031
;; INFORMATION FOR SEQ ID NO: 3:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 108 amino acids
;; TYPE: amino acid
;; STRANDEDNESS: single
;; TOPOLOGY: linear
US-08-663-809-3

Query Match 67.4%; Score 31; DB 1; Length 108;
Best Local Similarity 75.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 LCTELQTT 8
Db 17 LCTELTYT 24

RESULT 37
US-08-049-503-1
; Sequence 1, Application US/08049503
; Patent No. 5494662
; GENERAL INFORMATION:
; APPLICANT: UENO, KOJI
; APPLICANT: KATAYAMA, TERUAKI
; APPLICANT: MIYAMOTO, TSUMORU
; TITLE OF INVENTION: STIMULATOR FOR BONE FORMATION
; NUMBER OF SEQUENCES: 12
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: STEVENS, DAVIS, MILLER & MOSHER, LLP
; STREET: 515 N. WASHINGTON ST.
; CITY: ALEXANDRIA
; STATE: VA
; COUNTRY: USA
; ZIP: 22314
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/049,503
; FILING DATE: 21-APR-1993
; CLASSIFICATION: 424
; ATTORNEY/AGENT INFORMATION:
; NAME: POULOS III, JAMES A.
; REGISTRATION NUMBER: 31,714
; REFERENCE/DOCKET NUMBER: TPP29045
; TELECOMMUNICATION INFORMATION:

TELEPHONE: 703-549-7200
TELEFAX: 703-528-5313
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
ORGANISM: Homo sapiens
US-08-049-503-1

Query Match 67.4%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 LCTELQTT 8
Db 23 LCTELTWT 30

RESULT 38
US-08-225-224-2
Sequence 2, Application US/08225224
Patent No. 5635599
GENERAL INFORMATION:
APPLICANT: PASTAN, Ira
APPLICANT: KREITMAN, Robert J.
TITLE OF INVENTION: CIRCULARLY PERMUTATED LIGANDS AND
TITLE OF INVENTION: CIRCULARLY PERMUTED FUSION PROTEINS
NUMBER OF SEQUENCES: 57
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend Kourie and Crew
STREET: Stewart Street Tower, One Market Plaza
CITY: San Francisco
STATE: California
COUNTRY: US
ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/225,224
FILING DATE: 8-APR-1994
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen L.
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 15280-193
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 543-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULAR TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..129
OTHER INFORMATION: /label= IL4
US-08-225-224-2

Query Match 67.4%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;

Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
Qy 1 LCTELQTT 8
Db 23 LCTELTWT 30

RESULT 39
US-08-470-299-21
Sequence 21, Application US/08470299
Patent No. 5783181
GENERAL INFORMATION:
APPLICANT: Browne, Michael J.
APPLICANT: Murphy, Kay E.
APPLICANT: Chapman, Conrad G.
APPLICANT: Clinkbeard, Helen E.
APPLICANT: Young, Peter R.
APPLICANT: Shatzman, Allan R.
TITLE OF INVENTION: No. 5783181el Compounds
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESSER: SmithKline Beecham Corporation
STREET: 709 Swedeland Road, P.O. Box 1539
CITY: King of Prussia
STATE: Pennsylvania
COUNTRY: USA
ZIP: 19406
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/470,299
FILING DATE: 06-JUN-1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sutton, Jeffrey A.
REGISTRATION NUMBER: 34,028
REFERENCE/DOCKET NUMBER: P31005C3
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5024
TELEFAX: 610-270-5090
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
US-08-470-299-21

Query Match 67.4%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 LCTELQTT 8
Db 23 LCTELTWT 30

RESULT 40
US-08-874-697-1
Sequence 1, Application US/08874697
Patent No. 5986059
GENERAL INFORMATION:
APPLICANT: Shanefelt, Armen; Greve, Jeffrey; Gundel, Robert
TITLE OF INVENTION: T-cell Selective Interleukin-4 Agonists
NUMBER OF SEQUENCES: 42
CORRESPONDENCE ADDRESS:

ADDRESSEE: Bayer Corporation, Pharmaceutical Division
STREET: 400 Morgan Lane
CITY: West Haven
STATE: CT
COUNTRY: United States of America
ZIP: 06516-4175
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS v. 6.30
SOFTWARE: Word for Windows 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/874,697
FILING DATE:
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/019,748
FILING DATE: 14-JUN-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/036,746
FILING DATE: 27-JAN-1997
ATTORNEY/AGENT INFORMATION:
NAME: Huw R. Jones
REGISTRATION NUMBER: 33, 916
REFERENCE/DOCKET NUMBER: WH5013
TELECOMMUNICATION INFORMATION:
TELEPHONE: (203) 812-2317
TELEFAX: (203) 812-5492
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 129
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
DESCRIPTION: human Interleukin-4 protein
HYPOTHETICAL: no
ANTI-SENSE: no
US-08-874-697-1

Query Match 67.4%; Score 31; DB 1; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 LCTELQTT 8
Db 23 LCTELVT 30

RESULT 41
US-08-722-258-2
Sequence 2, Application US/08722258
Patent No. 6011002
GENERAL INFORMATION:
APPLICANT: Pastan, Ira
APPLICANT: Kreitman, Robert J.
APPLICANT: Furt, Raj K.
TITLE OF INVENTION: Circularly Permuted Ligands and
TITLE OF INVENTION: Circularly Permuted Chimeric Molecules
NUMBER OF SEQUENCES: 72
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: California
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/722,258
FILING DATE: 08-JAN-1997
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/US95/04468
FILING DATE: 06-APR-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/225,224
FILING DATE: 08-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 015280-193100US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
STRANDEDNESS:
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..129
OTHER INFORMATION: /note= "interleukin 4 (IL4)"
US-08-722-258-2

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Oy 1 LCTELQTT 8
Db 23 LCTELVT 30

RESULT 42
US-08-897-020-1
Sequence 1, Application US/08897020
Patent No. 6028176
GENERAL INFORMATION:
APPLICANT: Shanafelt, Armen; Greve, Jeffrey; Rocznik, Steven
TITLE OF INVENTION: High-affinity Interleukin-4 Mucins
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bayer Corporation, Pharmaceutical Division
STREET: 400 Morgan Lane
CITY: West Haven
STATE: CT
COUNTRY: United States of America
ZIP: 06516-4175
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS v. 6.30
SOFTWARE: Word for Windows 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/897,020
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: P-91,242
FILING DATE: 19-JUL-1996
ATTORNEY/AGENT INFORMATION:
NAME: Huw R. Jones
REGISTRATION NUMBER: 33, 916
REFERENCE/DOCKET NUMBER: WH5020
TELECOMMUNICATION INFORMATION:
TELEPHONE: (203) 812-2317
TELEFAX: (203) 812-5492
INFORMATION FOR SEQ ID NO: 1:

SEQUENCE CHARACTERISTICS:
LENGTH: 129
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
DESCRIPTION: human Interleukin-4 protein
HYPOTHETICAL: no
ANTI-SENSE: no
US-08-897-020-1

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
DB 23 LCTELTWT 30

RESULT 43
US-08-765-012A-16
Sequence 16, Application US/08765012A
Patent No. 6130318
GENERAL INFORMATION:
APPLICANT: Wild, Hanno; Hanks, Rudolf; Dorschung, Michael;
APPLICANT: Horlein, Hans-Dietrich; Beumink, Jürgen;
APPLICANT: Apeler, Heiner; Wehlmann, Hermann; and Sebald,
APPLICANT: Walter
TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS
TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sprung Kramer Schaefer & Briscoe
STREET: 660 White Plains Road
CITY: Tarrytown
STATE: New York
COUNTRY: USA
ZIP: 10591-5144
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
COMPUTER: Apple Macintosh
OPERATING SYSTEM: System 7.5
SOFTWARE: WordPerfect 3.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,012A
FILING DATE: 19-DEC-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/02358
FILING DATE: 19-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 44 23 131
FILING DATE: 01-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
REGISTRATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 332-1700
TELEFAX: (914) 332-1844
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
US-08-765-012A-16

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
DB 23 LCTELTWT 30

RESULT 44
US-08-765-012A-17
Sequence 17, Application US/08765012A
Patent No. 6130318
GENERAL INFORMATION:
APPLICANT: Wild, Hanno; Hanks, Rudolf; Dorschung, Michael;
APPLICANT: Horlein, Hans-Dietrich; Beumink, Jürgen;
APPLICANT: Apeler, Heiner; Wehlmann, Hermann; and Sebald,
APPLICANT: Walter
TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS
TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sprung Kramer Schaefer & Briscoe
STREET: 660 White Plains Road
CITY: Tarrytown
STATE: New York
COUNTRY: USA
ZIP: 10591-5144
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
COMPUTER: Apple Macintosh
OPERATING SYSTEM: System 7.5
SOFTWARE: WordPerfect 3.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/765,012A
FILING DATE: 19-DEC-1996
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/EP95/02358
FILING DATE: 19-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE 44 23 131
FILING DATE: 01-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Kurt G. Briscoe
REGISTRATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 332-1700
TELEFAX: (914) 332-1844
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
US-08-765-012A-17

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
DB 23 LCTELTWT 30

RESULT 45
US-08-765-012A-18
Sequence 18, Application US/08765012A
Patent No. 6130318
GENERAL INFORMATION:
APPLICANT: Wild, Hanno; Hanks, Rudolf; Dorschung, Michael;
APPLICANT: Horlein, Hans-Dietrich; Beumink, Jürgen;
APPLICANT: Apeler, Heiner; Wehlmann, Hermann; and Sebald,
APPLICANT: Walter
TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS

;; TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
;; TITLE OF INVENTION: HUMAN INTERLEUKIN 4
;; NUMBER OF SEQUENCES: 20
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Sprung Kramer Schaefer & Briscoe
;; STREET: 660 White Plains Road
;; CITY: Tarrytown
;; STATE: New York
;; COUNTRY: USA
;; ZIP: 10591-5144
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
;; COMPUTER: Apple Macintosh
;; OPERATING SYSTEM: System 7.5
;; SOFTWARE: WordPerfect 3.5
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/765, 012A
;; FILING DATE: 19-DEC-1996
;; CLASSIFICATION: 435
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: PCT/EP95/02358
;; FILING DATE: 19-JUN-1995
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: DE 44 23 131
;; FILING DATE: 01-JUL-1994
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Kurt G. Briscoe
;; REGISTRATION NUMBER: 33,141
;; REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (914) 332-1700
;; TELEFAX: (914) 332-1844
;; INFORMATION FOR SEQ ID NO: 18:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 129 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; US-08-765-012A-18
;;
Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
QY 1 LCTELQTT 8
Db 23 LCTELVT 30
;;
RESULT 46
;; US-08-765-012A-19
;; Sequence 19, Application US/08765012A
;; Patent No. 6130318
;; GENERAL INFORMATION:
;; APPLICANT: Wild, Hanno; Hanko, Rudolf; Dorschug, Michael;
;; APPLICANT: Horlein, Hans-Dietrich; Beunink, Jürgen;
;; APPLICANT: Apelzer, Heiner; Wehlmann, Hermann; and Sebald,
;; APPLICANT: Walter
;; TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS
;; TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
;; TITLE OF INVENTION: HUMAN INTERLEUKIN 4
;; NUMBER OF SEQUENCES: 20
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Sprung Kramer Schaefer & Briscoe
;; STREET: 660 White Plains Road
;; CITY: Tarrytown
;; STATE: New York
;; COUNTRY: USA
;; ZIP: 10591-5144
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
;; COMPUTER: Apple Macintosh
;; OPERATING SYSTEM: System 7.5
;; SOFTWARE: WordPerfect 3.5

;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/765, 012A
;; FILING DATE: 19-DEC-1996
;; CLASSIFICATION: 435
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: PCT/EP95/02358
;; FILING DATE: 19-JUN-1995
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: DE 44 23 131
;; FILING DATE: 01-JUL-1994
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Kurt G. Briscoe
;; REGISTRATION NUMBER: 33,141
;; REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
;; TELECOMMUNICATION INFORMATION:
;; TELEPHONE: (914) 332-1700
;; TELEFAX: (914) 332-1844
;; INFORMATION FOR SEQ ID NO: 19:
;; SEQUENCE CHARACTERISTICS:
;; LENGTH: 129 amino acids
;; TYPE: amino acid
;; TOPOLOGY: linear
;; US-08-765-012A-19
;;
Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
QY 1 LCTELQTT 8
Db 23 LCTELVT 30
;;
RESULT 47
;; US-08-765-012A-20
;; Sequence 20, Application US/08765012A
;; Patent No. 6130318
;; GENERAL INFORMATION:
;; APPLICANT: Wild, Hanno; Hanko, Rudolf; Dorschug, Michael;
;; APPLICANT: Horlein, Hans-Dietrich; Beunink, Jürgen;
;; APPLICANT: Apelzer, Heiner; Wehlmann, Hermann; and Sebald,
;; APPLICANT: Walter
;; TITLE OF INVENTION: HIL-4 MUTANT PROTEINS USED AS
;; TITLE OF INVENTION: ANTAGONISTS OR PARTIAL AGONISTS OF
;; TITLE OF INVENTION: HUMAN INTERLEUKIN 4
;; NUMBER OF SEQUENCES: 20
;; CORRESPONDENCE ADDRESS:
;; ADDRESSEE: Sprung Kramer Schaefer & Briscoe
;; STREET: 660 White Plains Road
;; CITY: Tarrytown
;; STATE: New York
;; COUNTRY: USA
;; ZIP: 10591-5144
;; COMPUTER READABLE FORM:
;; MEDIUM TYPE: Diskette, 3.50 inch, 1.4 Mb storage
;; COMPUTER: Apple Macintosh
;; OPERATING SYSTEM: System 7.5
;; SOFTWARE: WordPerfect 3.5
;; CURRENT APPLICATION DATA:
;; APPLICATION NUMBER: US/08/765, 012A
;; FILING DATE: 19-DEC-1996
;; CLASSIFICATION: 435
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: PCT/EP95/02358
;; FILING DATE: 19-JUN-1995
;; PRIOR APPLICATION DATA:
;; APPLICATION NUMBER: DE 44 23 131
;; FILING DATE: 01-JUL-1994
;; ATTORNEY/AGENT INFORMATION:
;; NAME: Kurt G. Briscoe
;; REGISTRATION NUMBER: 33,141
;; REFERENCE/DOCKET NUMBER: BAYER 9776-KGB
;; TELECOMMUNICATION INFORMATION:

TELEPHONE: (914) 332-1700
TELEFAX: (914) 332-1844
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 129 amino acids
TYPE: amino acid
TOPOLOGY: linear
US-08-765-012A-20

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
| | | | |
Db 23 LCTELTVT 30

RESULT 48

US-09-350-823-1
Sequence 1, Application US/09350823
Patent No. 6313272

GENERAL INFORMATION:
APPLICANT: Shanafelt, Armen; Greve, Jeffrey; Rocznik, Steven
TITLE OF INVENTION: High-affinity Interleukin-4 Mutains
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSER: Bayer Corporation, Pharmaceutical Division
STREET: 400 Morgan Lane
CITY: West Haven
STATE: CT
COUNTRY: United States of America
ZIP: 06516-4175

COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb storage
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS V. 6.30
SOFTWARE: Word for Windows 6.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/350,823
FILING DATE: 09-Jul-1999
CLASSIFICATION: <Unknown>

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/897,020
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Huw R. Jones
REGISTRATION NUMBER: 33, 916
REFERENCE/DOCKET NUMBER: WH5020

TELECOMMUNICATION INFORMATION:
TELEPHONE: (203) 812-2317
TELEFAX: (203) 812-5492
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:

LENGTH: 129
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
DESCRIPTION: human Interleukin-4 protein
HYPOTHETICAL: no
ANTI-SENSE: no
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-350-823-1

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
| | | | |
Db 23 LCTELTVT 30

RESULT 49
US-09-298-374-1
Sequence 1, Application US/09298374
Patent No. 6335426

GENERAL INFORMATION:
APPLICANT: Shanafelt, Armen B.
APPLICANT: Greve, Jeffrey M.
APPLICANT: Gundel, Robert

TITLE OF INVENTION: T-CELL SELECTIVE INTERLEUKIN-4 AGONISTS
FILE REFERENCE: 5013P2
CURRENT APPLICATION NUMBER: US/09/298,374
CURRENT FILING DATE: 1999-04-23

PRIOR APPLICATION NUMBER: 08/874,697
PRIOR FILING DATE: 1997-06-13
PRIOR APPLICATION NUMBER: 60/036,746
PRIOR FILING DATE: 1997-01-27
PRIOR APPLICATION NUMBER: 60/019,748
PRIOR FILING DATE: 1996-06-14

NUMBER OF SEQ ID NOS: 42
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO: 1
LENGTH: 129

TYPE: PRT
ORGANISM: Homo sapiens
FEATURES:
US-09-298-374-1

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
| | | | |
Db 23 LCTELTVT 30

RESULT 50
US-08-271-539-46
Sequence 46, Application US/08271539
Patent No. 6358509

GENERAL INFORMATION:
APPLICANT: Ramanathan, Lata
APPLICANT: Seelig, Gail F.
APPLICANT: Troceta, Paul P.

TITLE OF INVENTION: Antibody Antagonists of Human Interleukin-4
FILE REFERENCE: JB0059KO US
CURRENT APPLICATION NUMBER: US/08/271,539
CURRENT FILING DATE: 1994-07-07

PRIOR APPLICATION NUMBER: US 07/453,570
PRIOR FILING DATE: 1989-12-20
PRIOR APPLICATION NUMBER: PCT/US90/07289
PRIOR FILING DATE: 1990-12-18
PRIOR APPLICATION NUMBER: US 07/859,689
PRIOR FILING DATE: 1992-06-11

NUMBER OF SEQ ID NOS: 46
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO: 46
LENGTH: 129

TYPE: PRT
ORGANISM: Homo sapiens
US-08-271-539-46

Query Match 67.4%; Score 31; DB 2; Length 129;
Best Local Similarity 75.0%; Pred. No. 1.7e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTT 8
| | | | |
Db 23 LCTELTVT 30

Search completed: May 5, 2006, 07:09:50

Job time : 29.75 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:50:57 ; Search time 57 Seconds
(without alignments)
65.973 Million cell updates/sec

Title: US-08-170-344-4
Perfect score: 46
Sequence: 1 LCTELQTTI 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Published Applications_AA_Main:
1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep.*
4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep.*
5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep.*
6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep.*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	46	100.0	15	4	US-10-476-570-20
2	46	100.0	15	4	US-10-476-570-21
3	46	100.0	21	4	US-10-476-570-8
4	46	100.0	25	6	US-11-021-949-1
5	46	100.0	30	4	US-10-476-570-53
6	46	100.0	30	5	US-10-858-384-4
7	46	100.0	32	4	US-10-476-570-9
8	46	100.0	33	4	US-10-476-570-19
9	46	100.0	151	5	US-10-484-063-27
10	46	100.0	151	5	US-10-484-063-20
11	46	100.0	151	5	US-10-858-384-2
12	46	100.0	158	5	US-10-367-057-16
13	46	100.0	158	6	US-11-021-949-13
14	46	100.0	158	6	US-10-472-724-2
15	46	100.0	171	4	US-11-072-288-1
16	46	100.0	243	6	US-09-367-309A-1
17	46	100.0	266	3	US-10-000-903-4
18	46	100.0	273	4	US-10-899-771-4
19	46	100.0	273	5	US-10-000-903-10
20	46	100.0	292	5	US-10-899-771-10
21	46	100.0	292	5	US-10-000-903-6
22	46	100.0	371	5	US-10-899-771-6
23	46	100.0	371	5	US-10-899-771-14
24	46	100.0	390	5	US-10-899-771-14
25	46	100.0	390	5	US-10-899-771-14
26	46	100.0	536	4	US-10-367-059-10
27	46	100.0	536	4	US-10-368-046-10

28	46	100.0	536	4	US-10-367-367-10	Sequence 10, Appl
29	46	100.0	536	5	US-10-918-337-10	Sequence 10, Appl
30	38	82.6	239	5	US-10-820-155-14	Sequence 14, Appl
31	38	82.6	239	5	US-10-820-155-23	Sequence 23, Appl
32	38	82.6	239	5	US-10-820-155-30	Sequence 30, Appl
33	38	82.6	239	5	US-10-820-155-39	Sequence 39, Appl
34	38	82.6	239	5	US-10-820-155-85	Sequence 85, Appl
35	38	82.6	239	5	US-10-820-155-94	Sequence 94, Appl
36	38	82.6	239	5	US-10-820-155-97	Sequence 97, Appl
37	37	80.4	48	3	US-09-925-299-1386	Sequence 1386, Ap
38	37	80.4	48	3	US-09-925-299-1386	Sequence 1386, Ap
39	37	80.4	158	4	US-10-282-122A-70249	Sequence 70249, A
40	37	80.4	158	5	US-10-857-625-567	Sequence 567, App
41	36	78.3	452	5	US-10-424-599-191823	Sequence 191823,
42	36	78.3	452	5	US-10-450-763-17868	Sequence 37868, A
43	36	78.3	759	4	US-10-408-765A-748	Sequence 748, App
44	35	76.1	25	6	US-11-021-949-2	Sequence 2, Appl1
45	35	76.1	25	6	US-11-021-949-57	Sequence 57, Appl
46	35	76.1	42	5	US-10-751-845-152	Sequence 152, App
47	35	76.1	82	4	US-10-437-963-175677	Sequence 175677,
48	35	76.1	119	5	US-10-751-845-159	Sequence 159, App
49	35	76.1	158	5	US-10-800-023-27	Sequence 27, Appl
50	35	76.1	158	6	US-11-021-949-28	Sequence 28, Appl
51	35	76.1	158	6	US-11-021-949-29	Sequence 29, Appl
52	35	76.1	172	4	US-10-472-724-6	Sequence 6, Appl1
53	35	76.1	236	5	US-10-751-845-157	Sequence 157, App
54	35	76.1	237	5	US-10-751-845-158	Sequence 158, App
55	35	76.1	261	5	US-10-751-845-160	Sequence 160, App
56	35	76.1	278	5	US-10-899-771-21	Sequence 21, Appl
57	35	76.1	278	5	US-10-899-771-21	Sequence 21, Appl
58	35	76.1	383	5	US-10-000-903-23	Sequence 23, Appl
59	35	76.1	383	5	US-10-899-771-23	Sequence 23, Appl
60	35	76.1	801	4	US-10-424-599-226123	Sequence 226123,
61	35	76.1	946	5	US-10-732-923-8378	Sequence 8378, Ap
62	34	73.9	95	4	US-10-425-115-212369	Sequence 212369,
63	34	73.9	120	4	US-10-767-701-37218	Sequence 37218, A
64	34	73.9	244	5	US-10-820-155-106	Sequence 106, App
65	34	73.9	587	5	US-10-369-493-1841	Sequence 1841, Ap
66	34	73.9	587	5	US-10-732-923-11310	Sequence 11310, A
67	34	73.9	596	5	US-10-732-923-11311	Sequence 11311, A
68	34	73.9	637	5	US-10-668-936-7	Sequence 936-7, A
69	33	71.7	15	4	US-10-476-570-22	Sequence 22, Appl
70	33	71.7	74	4	US-10-425-115-292434	Sequence 292434,
71	33	71.7	82	4	US-10-425-115-179241	Sequence 179241,
72	33	71.7	137	4	US-10-425-115-189580	Sequence 189580,
73	33	71.7	138	4	US-10-425-115-33528	Sequence 33528,
74	33	71.7	158	6	US-11-021-949-30	Sequence 30, Appl
75	33	71.7	158	6	US-11-021-949-361	Sequence 361, App
76	33	71.7	179	4	US-10-424-599-147599	Sequence 147599,
77	33	71.7	212	4	US-10-363-616-453	Sequence 453, App
78	33	71.7	227	3	US-09-925-302-604	Sequence 604, App
79	33	71.7	227	3	US-09-925-302-604	Sequence 604, App
80	33	71.7	252	4	US-10-767-701-43809	Sequence 43809, A
81	33	71.7	252	5	US-10-739-930-6393	Sequence 6393, Ap
82	33	71.7	312	4	US-10-424-599-147748	Sequence 147748,
83	33	71.7	329	4	US-10-424-599-147748	Sequence 147707,
84	33	71.7	332	4	US-10-282-122A-69154	Sequence 69154, A
85	33	71.7	342	4	US-10-425-114-47110	Sequence 47110, A
86	33	71.7	392	5	US-10-450-763-57233	Sequence 57233, A
87	33	71.7	395	5	US-10-954-778-41	Sequence 41, Appl
88	33	71.7	574	4	US-10-437-963-102998	Sequence 102998,
89	33	71.7	662	6	US-11-097-143-12708	Sequence 12708, A
90	33	71.7	863	4	US-10-437-963-184291	Sequence 184291, A
91	32	69.6	56	4	US-10-083-357-960	Sequence 960, App
92	32	69.6	56	4	US-10-424-599-264561	Sequence 264561,
93	32	69.6	76	3	US-09-867-550-1112	Sequence 1112, Ap
94	32	69.6	78	4	US-10-424-599-200143	Sequence 200143,
95	32	69.6	87	4	US-10-425-115-24195	Sequence 24195,
96	32	69.6	101	4	US-10-767-701-54467	Sequence 54467, A
97	32	69.6	104	4	US-10-424-599-246277	Sequence 246277,
98	32	69.6	114	6	US-11-097-143-36090	Sequence 36090, A
99	32	69.6	147	4	US-10-443-201-64	Sequence 64, Appl
100	32	69.6	147	4	US-10-443-201-66	Sequence 66, Appl

101	32	69.6	147	4	US-10-443-201-68	Sequence 68, Appl	174	31	67.4	130	4	US-10-053-155A-4	Sequence 4, Appl1
102	32	69.6	147	4	US-10-443-201-70	Sequence 70, Appl	175	31	67.4	130	5	US-10-820-559-9	Sequence 9, Appl1
103	32	69.6	149	4	US-10-425-115-229668	Sequence 229668,	176	31	67.4	130	5	US-10-820-559-11	Sequence 11, Appl
104	32	69.6	180	4	US-10-425-115-187766	Sequence 187766,	177	31	67.4	130	5	US-10-820-559-12	Sequence 12, Appl
105	32	69.6	183	4	US-10-425-115-207590	Sequence 207590,	178	31	67.4	130	5	US-10-820-559-13	Sequence 13, Appl
106	32	69.6	341	5	US-10-450-763-54568	Sequence 54568, A	179	31	67.4	130	5	US-10-820-559-14	Sequence 14, Appl
107	32	69.6	398	5	US-10-954-778-39	Sequence 39, Appl	180	31	67.4	130	5	US-10-820-559-15	Sequence 15, Appl
108	32	69.6	472	4	US-10-424-559-165549	Sequence 165549,	181	31	67.4	130	5	US-10-820-559-16	Sequence 16, Appl
109	32	69.6	555	4	US-10-767-701-46075	Sequence 46075, A	182	31	67.4	134	4	US-10-437-963-112241	Sequence 112241,
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111	32	69.6	613	4	US-10-425-114-63762	Sequence 63762, A	184	31	67.4	149	6	US-11-021-949-16	Sequence 16, Appl
112	32	69.6	613	4	US-10-425-115-343502	Sequence 343502,	185	31	67.4	150	4	US-10-233-902-25	Sequence 25, Appl
113	32	69.6	618	6	US-11-097-143-25152	Sequence 25152, A	186	31	67.4	150	6	US-11-041-636-25	Sequence 25, Appl
114	32	69.6	684	4	US-10-437-963-184182	Sequence 184182,	187	31	67.4	153	3	US-09-947-770-13	Sequence 13, Appl
115	32	69.6	853	4	US-10-437-963-122232	Sequence 122232,	188	31	67.4	153	3	US-09-923-246-112	Sequence 112, App
116	32	69.6	1889	4	US-10-437-963-184639	Sequence 184639,	189	31	67.4	153	4	US-10-235-123-112	Sequence 112, App
117	31	67.4	42	4	US-10-425-115-283541	Sequence 283541,	190	31	67.4	153	4	US-10-282-522-8	Sequence 8, Appl1
118	31	67.4	60	4	US-10-425-115-318928	Sequence 318928,	191	31	67.4	153	4	US-10-351-157-179	Sequence 179, App
119	31	67.4	63	4	US-10-437-963-202918	Sequence 202918,	192	31	67.4	153	4	US-10-352-554-164	Sequence 164, App
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122	31	67.4	85	4	US-10-437-963-156344	Sequence 156344,	195	31	67.4	153	4	US-10-683-516-1	Sequence 1, Appl1
123	31	67.4	107	4	US-10-437-963-165626	Sequence 165626,	196	31	67.4	153	5	US-10-688-845-6	Sequence 6, Appl1
124	31	67.4	108	5	US-10-450-763-32015	Sequence 32015, A	197	31	67.4	153	5	US-10-688-845-6	Sequence 6, Appl1
125	31	67.4	109	5	US-10-450-763-46825	Sequence 46825, A	198	31	67.4	153	5	US-10-787-442-112	Sequence 112, App
126	31	67.4	110	4	US-10-424-559-144591	Sequence 144591,	199	31	67.4	153	5	US-10-951-239-6	Sequence 6, Appl1
127	31	67.4	128	4	US-10-150-874-2	Sequence 2, Appl1	200	31	67.4	153	5	US-10-775-304-178	Sequence 178, App
128	31	67.4	129	3	US-09-792-793A-25	Sequence 25, Appl1	201	31	67.4	187	5	US-10-450-763-4764	Sequence 47624, A
129	31	67.4	129	3	US-09-969-748C-27	Sequence 27, Appl1	202	31	67.4	195	5	US-10-450-763-3525	Sequence 3525, A
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131	31	67.4	129	4	US-10-050-227-21	Sequence 21, Appl1	204	31	67.4	203	4	US-10-425-114-7115	Sequence 71115, A
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133	31	67.4	129	4	US-10-400-708-11	Sequence 11, Appl	206	31	67.4	226	5	US-10-450-763-37872	Sequence 37872, A
134	31	67.4	129	4	US-10-298-148-11	Sequence 11, Appl	207	31	67.4	226	5	US-10-450-763-35338	Sequence 35338, A
135	31	67.4	129	4	US-10-375-209A-25	Sequence 25, Appl1	208	31	67.4	230	5	US-10-450-763-46139	Sequence 46139, A
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137	31	67.4	129	4	US-10-658-834A-207	Sequence 207, App	210	31	67.4	256	4	US-10-437-963-133260	Sequence 133260,
138	31	67.4	129	4	US-10-658-834A-546	Sequence 546, App	211	31	67.4	256	4	US-10-424-599-158115	Sequence 158115,
139	31	67.4	129	4	US-10-658-834A-547	Sequence 547, App	212	31	67.4	260	5	US-10-450-763-38197	Sequence 38197, A
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142	31	67.4	129	4	US-10-658-834A-550	Sequence 550, App	215	31	67.4	277	5	US-10-450-763-40580	Sequence 40580, A
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146	31	67.4	129	4	US-10-658-834A-554	Sequence 554, App	219	31	67.4	295	5	US-10-450-763-36622	Sequence 36622, A
147	31	67.4	129	4	US-10-658-834A-555	Sequence 555, App	220	31	67.4	295	5	US-10-450-763-58180	Sequence 58140, A
148	31	67.4	129	4	US-10-658-834A-556	Sequence 556, App	221	31	67.4	300	5	US-10-450-763-53559	Sequence 53559, A
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153	31	67.4	129	4	US-10-658-834A-561	Sequence 561, App	226	31	67.4	327	5	US-10-450-763-35246	Sequence 35246, A
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161	31	67.4	129	4	US-10-773-939-11	Sequence 11, Appl	234	31	67.4	342	3	US-09-804-291-207	Sequence 207, App
162	31	67.4	129	4	US-10-773-939-11	Sequence 11, Appl	235	31	67.4	342	5	US-10-450-763-47690	Sequence 47690, A
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165	31	67.4	129	5	US-10-872-198-118	Sequence 118, App	238	31	67.4	348	5	US-10-450-763-39215	Sequence 39215, A
166	31	67.4	129	5	US-10-872-198-118	Sequence 118, App	239	31	67.4	352	5	US-10-450-763-58440	Sequence 58440, A
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263	31	67.4	401	5	US-10-450-763-44631	Sequence 44631, A	336	31	67.4	520	5	US-10-450-763-34219	Sequence 34219, A
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869	31	67.4	1104	5	US-10-450-763-34701	Sequence 34701, A	942	31	67.4	1143	5	US-10-450-763-36980	Sequence 47851, A
870	31	67.4	1104	5	US-10-450-763-36431	Sequence 36431, A	943	31	67.4	1144	5	US-10-450-763-53562	Sequence 53562, A
871	31	67.4	1104	5	US-10-450-763-36976	Sequence 36976, A	944	31	67.4	1145	5	US-10-450-763-37782	Sequence 37782, A
872	31	67.4	1104	5	US-10-450-763-41172	Sequence 41172, A	945	31	67.4	1150	5	US-10-450-763-44308	Sequence 44308, A
873	31	67.4	1105	5	US-10-450-763-40491	Sequence 40491, A	946	31	67.4	1150	5	US-10-450-763-45129	Sequence 45129, A
874	31	67.4	1105	5	US-10-450-763-35543	Sequence 35543, A	947	31	67.4	1150	5	US-10-450-763-50235	Sequence 50235, A
875	31	67.4	1106	5	US-10-450-763-40447	Sequence 40447, A	948	31	67.4	1152	5	US-10-450-763-36432	Sequence 36432, A
876	31	67.4	1108	5	US-10-450-763-44951	Sequence 44951, A	949	31	67.4	1152	5	US-10-450-763-40013	Sequence 40013, A
877	31	67.4	1108	5	US-10-450-763-45264	Sequence 45264, A	950	31	67.4	1152	5	US-10-450-763-32542	Sequence 32542, A
878	31	67.4	1108	5	US-10-450-763-47548	Sequence 47548, A	951	31	67.4	1153	5	US-10-450-763-43357	Sequence 34357, A
879	31	67.4	1108	5	US-10-450-763-48818	Sequence 48818, A	952	31	67.4	1153	5	US-10-450-763-47850	Sequence 47850, A
880	31	67.4	1108	5	US-10-450-763-53529	Sequence 53529, A	953	31	67.4	1153	5	US-10-450-763-39500	Sequence 39500, A
881	31	67.4	1110	5	US-10-450-763-33869	Sequence 33869, A	954	31	67.4	1156	5	US-10-450-763-37418	Sequence 37418, A
882	31	67.4	1110	5	US-10-450-763-36986	Sequence 36986, A	955	31	67.4	1156	5	US-10-450-763-34344	Sequence 34344, A
883	31	67.4	1111	5	US-10-450-763-36440	Sequence 36440, A	956	31	67.4	1159	5	US-10-450-763-36975	Sequence 36975, A
884	31	67.4	1111	5	US-10-450-763-36983	Sequence 36983, A	957	31	67.4	1159	5	US-10-450-763-35994	Sequence 35994, A
885	31	67.4	1111	5	US-10-450-763-40020	Sequence 40020, A	958	31	67.4	1159	5	US-10-450-763-39474	Sequence 39474, A
886	31	67.4	1111	5	US-10-450-763-41180	Sequence 41180, A	959	31	67.4	1159	5	US-10-450-763-47980	Sequence 47980, A
887	31	67.4	1111	5	US-10-450-763-44956	Sequence 44956, A	960	31	67.4	1159	5	US-10-450-763-34769	Sequence 34769, A
888	31	67.4	1111	5	US-10-450-763-45127	Sequence 45127, A	961	31	67.4	1161	5	US-10-450-763-37343	Sequence 37343, A
889	31	67.4	1111	5	US-10-450-763-53538	Sequence 53538, A	962	31	67.4	1167	5	US-10-450-763-34344	Sequence 34344, A
890	31	67.4	1115	5	US-10-450-763-36436	Sequence 36436, A	963	31	67.4	1170	5	US-10-450-763-36975	Sequence 36975, A
891	31	67.4	1115	5	US-10-450-763-36443	Sequence 36443, A	964	31	67.4	1170	5	US-10-450-763-39006	Sequence 39006, A
892	31	67.4	1115	5	US-10-450-763-40016	Sequence 40016, A	965	31	67.4	1170	5	US-10-450-763-32006	Sequence 37418, A
893	31	67.4	1115	5	US-10-450-763-40023	Sequence 40023, A	966	31	67.4	1171	5	US-10-450-763-37418	Sequence 37418, A
894	31	67.4	1115	5	US-10-450-763-41176	Sequence 41176, A	967	31	67.4	1171	5	US-10-450-763-36972	Sequence 36972, A
895	31	67.4	1115	5	US-10-450-763-41183	Sequence 41183, A	968	31	67.4	1172	5	US-10-450-763-41366	Sequence 41366, A
896	31	67.4	1115	5	US-10-450-763-43637	Sequence 43637, A	969	31	67.4	1175	5	US-10-450-763-32695	Sequence 32695, A
897	31	67.4	1115	5	US-10-450-763-43637	Sequence 43637, A	970	31	67.4	1177	5	US-10-450-763-58550	Sequence 58550, A
898	31	67.4	1115	5	US-10-450-763-45128	Sequence 45128, A	971	31	67.4	1177	5	US-10-450-763-41546	Sequence 41546, A
899	31	67.4	1115	5	US-10-450-763-45292	Sequence 45292, A	972	31	67.4	1181	5	US-10-450-763-47869	Sequence 47869, A
900	31	67.4	1115	5	US-10-450-763-47558	Sequence 47558, A	973	31	67.4	1183	5	US-10-450-763-46162	Sequence 46162, A
901	31	67.4	1115	5	US-10-450-763-47849	Sequence 47849, A	974	31	67.4	1183	5	US-10-450-763-53544	Sequence 53544, A
902	31	67.4	1115	5	US-10-450-763-48816	Sequence 48816, A	975	31	67.4	1183	5	US-10-450-763-32697	Sequence 32697, A
903	31	67.4	1115	5	US-10-450-763-53551	Sequence 53551, A	976	31	67.4	1184	5	US-10-450-763-32697	Sequence 32697, A

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977 31 67.4 1185 5 US-10-450-763-39035 Sequence 39035, A
978 31 67.4 1186 5 US-10-450-763-39490 Sequence 39490, A
979 31 67.4 1189 5 US-10-450-763-32525 Sequence 32525, A
980 31 67.4 1189 5 US-10-450-763-34340 Sequence 34340, A
981 31 67.4 1189 5 US-10-450-763-36962 Sequence 36962, A
982 31 67.4 1189 5 US-10-450-763-38200 Sequence 38200, A
983 31 67.4 1189 5 US-10-450-763-45101 Sequence 45101, A
984 31 67.4 1189 5 US-10-450-763-47834 Sequence 47834, A
985 31 67.4 1190 5 US-10-450-763-36001 Sequence 36001, A
986 31 67.4 1190 5 US-10-450-763-36978 Sequence 36978, A
987 31 67.4 1190 5 US-10-450-763-38253 Sequence 38253, A
988 31 67.4 1190 5 US-10-450-763-54029 Sequence 54029, A
989 31 67.4 1192 5 US-10-450-763-36424 Sequence 36424, A
990 31 67.4 1192 5 US-10-450-763-40005 Sequence 40005, A
991 31 67.4 1194 5 US-10-450-763-34364 Sequence 34364, A
992 31 67.4 1196 5 US-10-450-763-44969 Sequence 44969, A
993 31 67.4 1206 5 US-10-450-763-48807 Sequence 48807, A
994 31 67.4 1206 5 US-10-450-763-53517 Sequence 53517, A
995 31 67.4 1210 5 US-10-450-763-47988 Sequence 47988, A
996 31 67.4 1212 5 US-10-450-763-34363 Sequence 34363, A
997 31 67.4 1213 5 US-10-450-763-48849 Sequence 48849, A
998 31 67.4 1213 5 US-10-450-763-53560 Sequence 53560, A
999 31 67.4 1214 5 US-10-450-763-39512 Sequence 39512, A
1000 31 67.4 1221 5 US-10-450-763-35253 Sequence 35253, A
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ALIGNMENTS

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RESULT 1
US-10-476-570-20
; Sequence 20, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 17-31
US-10-476-570-20

Query Match 100.0%; Score 46; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.14; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
|||
Db 6 LCTELQTTI 14
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RESULT 2
US-10-476-570-21
; Sequence 21, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
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APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 21
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 20-34
US-10-476-570-21
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Query Match 100.0%; Score 46; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.14; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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OY 1 LCTELQTTI 9
|||
Db 3 LCTELQTTI 11
```

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RESULT 3
US-10-476-570-8
; Sequence 8, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 8
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 14-34
US-10-476-570-8

Query Match 100.0%; Score 46; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.2; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
|||
Db 9 LCTELQTTI 17
```


RESULT 4
US-11-021-949-1
; Sequence 1, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SABIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-1

Query Match 100.0%; Score 46; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.24;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
DB 14 LCTELQTTI 22

RESULT 5
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIER, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53

Query Match 100.0%; Score 46; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.28;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
DB 8 LCTELQTTI 16

RESULT 6
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCES
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match 100.0%; Score 46; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.28;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
DB 8 LCTELQTTI 16

RESULT 7
US-10-476-570-9
; Sequence 9, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIER, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 32
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9

Query Match 100.0%; Score 46; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9

Db 9 LCTELQTTI 17

RESULT 8
US-10-476-570-19
; Sequence 19, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Geard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 33
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19

Query Match 100.0%; Score 46; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.31;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 9 LCTELQTTI 17

RESULT 9
US-10-177-390-6
; Sequence 6, Application US/10177390
; Publication No. US20030143743A1
; GENERAL INFORMATION:
; APPLICANT: Schuler, Gerold
; APPLICANT: N.V. Antwerp Innovatiecentrum
; TITLE OF INVENTION: Improved Transfection of Eucaryotic Cells with linear
; FILE REFERENCE: 021505wo/JH/ml
; CURRENT APPLICATION NUMBER: US/10/177,390
; CURRENT FILING DATE: 2002-06-20
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match 100.0%; Score 46; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 10

US-10-484-063-20
; Sequence 20, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILLEMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-20

Query Match 100.0%; Score 46; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 11
US-10-484-063-27
; Sequence 27, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILLEMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 27
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match 100.0%; Score 46; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 12
US-10-858-384-2
; Sequence 2, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE

APPLICANT: BOUGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCES
APPLICANT: FERRIS, ESTELLE
TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858,384
PRIORITY FILING DATE: 2004-06-02
PRIORITY FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match 100.0%; Score 46; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
|||||
DB 22 LCTELQTTI 30

RESULT 13
US-10-367-057-16
Sequence 16, Application US/10367057
Publication No. US20050100554A1
GENERAL INFORMATION:
APPLICANT: Cuthill, Scott;
APPLICANT: Jackson, Amanda;
APPLICANT: Lewin, David A.;
APPLICANT: Ooi, Chean Eng
TITLE OF INVENTION: Complexes and Methods of Using Same
FILE REFERENCE: 21402-559
CURRENT APPLICATION NUMBER: US/10/367,057
CURRENT FILING DATE: 2003-02-14
PRIORITY FILING DATE: 2002-02-14
PRIORITY FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 198
SOFTWARE: Curoseqblast version 0.1
SEQ ID NO 16
LENGTH: 158
TYPE: PRT
ORGANISM: Homo sapiens
US-10-367-057-16

Query Match 100.0%; Score 46; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
|||||
DB 22 LCTELQTTI 30

RESULT 14
US-11-021-949-13
Sequence 13, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
TITLE OF INVENTION: AND METHODS OF THEIR USE

FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIORITY FILING DATE: 2003-12-23
PRIORITY FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: PasteSeq for Windows Version 4.0
SEQ ID NO 13
LENGTH: 158
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match 100.0%; Score 46; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
|||||
DB 22 LCTELQTTI 30

RESULT 15
US-10-472-724-2
Sequence 2, Application US/10472724
Publication No. US20040171806A1
GENERAL INFORMATION:
APPLICANT: Cid-Arregui, Angel
APPLICANT: Zur Hausen, Harald
TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
FILE REFERENCE: 4121-154
CURRENT APPLICATION NUMBER: US/10/472,724
CURRENT FILING DATE: 2003-08-17
PRIORITY FILING DATE: PCT/EP02/03271
PRIORITY FILING DATE: 2002-03-22
PRIORITY FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.2
SEQ ID NO 2
LENGTH: 171
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match 100.0%; Score 46; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 1.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
|||||
DB 27 LCTELQTTI 35

RESULT 16
US-11-072-288-1
Sequence 1, Application US/11072288
Publication No. US20050159386A1
GENERAL INFORMATION:
APPLICANT: KIENY, Marie-Paule
APPLICANT: BALLOU, Jean-Marc
APPLICANT: BIZOUARNE, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
FILE REFERENCE: 01753-122
CURRENT APPLICATION NUMBER: US/11/072,288
CURRENT FILING DATE: 2005-03-07
PRIORITY FILING DATE: US/09/462,993
PRIORITY FILING DATE: 2000-04-17
PRIORITY FILING DATE: PCT/FR98/01576
PRIORITY FILING DATE: 1998-07-17

PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human Papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TWf.
US-11-072-288-1

Query Match 100.0%; Score 46; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 2.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
Db 50 LCTELQTTI 58

RESULT 17
US-09-367-309A-1
Sequence 1, Application US/09367309A
Publication No. US20020081329A1
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 46; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
Db 22 LCTELQTTI 30

RESULT 18
US-10-000-903-4
Sequence 4, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernandez
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22

NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-4

Query Match 100.0%; Score 46; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
Db 128 LCTELQTTI 136

RESULT 19
US-10-899-771-4
Sequence 4, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
OTHER INFORMATION: Influenzae B and E6 from Human papilloma virus type
OTHER INFORMATION: 16)
US-10-899-771-4

Query Match 100.0%; Score 46; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 2.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LCTELQTTI 9
Db 128 LCTELQTTI 136

RESULT 20
US-10-000-903-10
Sequence 10, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernandez
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5

PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-10

Query Match 100.0%; Score 46; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 2.7;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 147 LCTELQTTI 155

RESULT 21
US-10-899-771-10
; Sequence 10, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyra from Streptococcus
; OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-10

Query Match 100.0%; Score 46; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 2.7;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 147 LCTELQTTI 155

RESULT 22
US-10-000-903-6
; Sequence 6, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Caberon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17

PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-6

Query Match 100.0%; Score 46; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 128 LCTELQTTI 136

RESULT 23
US-10-899-771-6
; Sequence 6, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and E67 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-6

Query Match 100.0%; Score 46; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 3.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 128 LCTELQTTI 136

RESULT 24
US-10-000-903-14
; Sequence 14, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Caberon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285

;; PRIOR FILING DATE: 1998-08-17
;; PRIOR APPLICATION NUMBER: GB 9717953.5
;; PRIOR FILING DATE: 1997-08-22
;; NUMBER OF SEQ ID NOS: 23
;; SOFTWARE: FastSeq for Windows Version 3.0
;; SEQ ID NO 14
;; LENGTH: 390
;; TYPE: PRT
;; ORGANISM: Homo sapien
US-10-000-903-14

Query Match 100.0%; Score 46; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 3.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 147 LCTELQTTI 155

RESULT 25
US-10-899-771-14
;; Sequence 14, Application US/10899771
;; Publication No. US20050031638A1
;; GENERAL INFORMATION:
;; APPLICANT: Dalemans, Wilfried L.J.
;; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
;; TITLE OF INVENTION: and Fusion Proteins Adjuncted with a Cpg Oligonucleotide
;; FILE REFERENCE: B45124
;; CURRENT APPLICATION NUMBER: US/10/899,771
;; PRIOR FILING DATE: 2004-07-27
;; PRIOR APPLICATION NUMBER: US/09/581,976
;; PRIOR FILING DATE: 2000-06-20
;; PRIOR APPLICATION NUMBER: PCT/EP98/08563
;; PRIOR FILING DATE: 1998-12-18
;; PRIOR APPLICATION NUMBER: GB 9727262.9
;; PRIOR FILING DATE: 1997-12-24
;; NUMBER OF SEQ ID NOS: 28
;; SOFTWARE: FastSeq for Windows Version 3.0
;; SEQ ID NO 14
;; LENGTH: 390
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURES:
;; OTHER INFORMATION: Chimeraic protein (Clyta from Streptococcus
;; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
;; OTHER INFORMATION: virus type 16)
US-10-899-771-14

Query Match 100.0%; Score 46; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 3.6;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 147 LCTELQTTI 155

RESULT 26
US-10-367-095-10
;; Sequence 10, Application US/10367095
;; Publication No. US20030228696A1
;; GENERAL INFORMATION:
;; APPLICANT: Robin A. Robinson
;; TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
;; FILE REFERENCE: 44149-1US1
;; CURRENT APPLICATION NUMBER: US/10/367,095
;; PRIOR FILING DATE: 2003-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14

;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,135
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURES:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-095-10

Query Match 100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 492 LCTELQTTI 500

RESULT 27
US-10-368-046-10
;; Sequence 10, Application US/10368046
;; Publication No. US20040065188A1
;; GENERAL INFORMATION:
;; APPLICANT: Robin A. Robinson
;; TITLE OF INVENTION: Method for Isolation and Purification of
;; TITLE OF INVENTION: Expressed Gene Products In Vitro
;; FILE REFERENCE: 44149-3US1
;; CURRENT APPLICATION NUMBER: US/10/368,046
;; PRIOR FILING DATE: 2003-02-15
;; PRIOR APPLICATION NUMBER: US 60/356,119
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,161
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,118
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,133
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,157
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,135
;; PRIOR FILING DATE: 2002-02-14
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536

```

; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-368-046-10

```

```

Query Match          100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY      1 LCTELQTTI 9
        |||||
Db      492 LCTELQTTI 500

```

```

RESULT 28
US-10-367-367-10
; Sequence 10, Application US/10367367
; Publication No. US20040121465A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson

```

```

; TITLE OF INVENTION: Optimization of Gene Sequences of
; FILE REFERENCE: 44149-2US1
; CURRENT APPLICATION NUMBER: US/10/367,367
; PRIOR FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-367-10

```

```

Query Match          100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY      1 LCTELQTTI 9
        |||||
Db      492 LCTELQTTI 500

```

```

RESULT 29
US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:

```

```

; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; FILE REFERENCE: 19065/2132
; CURRENT APPLICATION NUMBER: US/10/918,337
; PRIOR FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161

```

```

; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10

```

```

Query Match          100.0%; Score 46; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY      1 LCTELQTTI 9
        |||||
Db      492 LCTELQTTI 500

```

```

RESULT 30
US-10-820-155-14
; Sequence 14, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:

```

```

; APPLICANT: Wellguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; PRIOR FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: Patencin version 3.1
; SEQ ID NO 14
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-14

```

```

Query Match          82.6%; Score 38; DB 5; Length 239;
Best Local Similarity 77.8%; Pred. No. 64;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 LCTELQTTI 9
        |||||
Db      145 LCTELQTTI 153

```

```

RESULT 31
US-10-820-155-23
; Sequence 23, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: Natimmune A/S
; APPLICANT: Wellguny, Dietmar

```


Qy 1 LCTELQTTI 9
| | | | |
Db 145 LCTELQTTV 153

RESULT 36
US-10-820-155-97
; Sequence 97, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: Natimmune A/S
; APPLICANT: Weiligun, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; CURRENT FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 97
; LENGTH: 239
; TYPE: PRT
; ORGANISM: Mus musculus
US-10-820-155-97

Query Match 82.6%; Score 38; DB 5; Length 239;
Best Local Similarity 77.8%; Pred. No. 64;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
| | | | |
Db 145 LCTELQTTV 153

RESULT 37
US-09-925-299-1386
; Sequence 1386, Application US/09925299
; Patent No. US20020055627A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA102
; CURRENT APPLICATION NUMBER: US/09/925,299
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05883
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 1556
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1386
; LENGTH: 48
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (5)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (13)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (15)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (40)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-299-1386

Query Match 80.4%; Score 37; DB 3; Length 48;

Best Local Similarity 87.5%; Pred. No. 20;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
Qy 1 LCTELQTT 8
| | | | |
Db 26 LCTELQTT 33

RESULT 38
US-09-925-299-1386
; Sequence 1386, Application US/09925299
; Publication No. US2003040617A9
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA102
; CURRENT APPLICATION NUMBER: US/09/925,299
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05883
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 1556
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 1386
; LENGTH: 48
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (5)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (13)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (15)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; NAME/KEY: SITE
; LOCATION: (40)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-09-925-299-1386

Query Match 80.4%; Score 37; DB 3; Length 48;
Best Local Similarity 87.5%; Pred. No. 20;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LCTELQTT 8
| | | | |
Db 26 LCTELQTT 33

RESULT 39
US-10-282-122A-70249
; Sequence 70249, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangshu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Haselbeck, Robert
; APPLICANT: Ohlsen, Karl
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Forsyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: ELPTRA 034A
; CURRENT APPLICATION NUMBER: US/10/282,122A
; CURRENT FILING DATE: 2003-02-20

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;; PRIOR APPLICATION NUMBER: 60/191,078
;; PRIOR FILING DATE: 2000-03-21
;; PRIOR APPLICATION NUMBER: 60/206,848
;; PRIOR FILING DATE: 2000-05-23
;; PRIOR APPLICATION NUMBER: 60/207,727
;; PRIOR FILING DATE: 2000-05-26
;; PRIOR APPLICATION NUMBER: 60/230,335
;; PRIOR FILING DATE: 2000-09-06
;; PRIOR APPLICATION NUMBER: 60/230,347
;; PRIOR FILING DATE: 2000-09-09
;; PRIOR APPLICATION NUMBER: 60/242,578
;; PRIOR FILING DATE: 2000-10-23
;; PRIOR APPLICATION NUMBER: 60/253,625
;; PRIOR FILING DATE: 2000-11-27/257,931
;; PRIOR APPLICATION NUMBER: 60/257,931
;; PRIOR FILING DATE: 2000-12-22
;; PRIOR APPLICATION NUMBER: 60/267,636
;; PRIOR FILING DATE: 2001-02-09
;; PRIOR APPLICATION NUMBER: 60/269,308
;; PRIOR FILING DATE: 2001-02-16
;; Remaining Prior Application data removed - See File Wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 78614
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO: 70249
;; LENGTH: 158
;; TYPE: PRT
;; ORGANISM: Staphylococcus aureus
US-10-282-122A-70249
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Query Match      80.4%; Score 37; DB 4; Length 158;
Best Local Similarity 66.7%; Pred. No. 64;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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```
Qy      1 LCTELQTTI 9
        |||:||||
Db      57 LCVEIQTTL 65
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RESULT 40
US-10-857-625-567
; Sequence 567, Application US/10857625
; Publication No. US20050026189A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Liangsu
; APPLICANT: Zamudio, Carlos
; TITLE OF INVENTION: MICROBIAL OPERONS
; FILE REFERENCE: ELITRA.036A
; CURRENT APPLICATION NUMBER: US/10/857,625
; CURRENT FILING DATE: 2004-05-28
; PRIOR APPLICATION NUMBER: 60/474768
; PRIOR FILING DATE: 2003-05-29
; NUMBER OF SEQ ID NOS: 833
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO: 567
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Staphylococcus aureus
US-10-857-625-567
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```
Query Match      80.4%; Score 37; DB 5; Length 158;
Best Local Similarity 66.7%; Pred. No. 64;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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```
Qy      1 LCTELQTTI 9
        |||:||||
Db      57 LCVEIQTTL 65
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RESULT 41
US-10-424-599-191823
; Sequence 191823, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
```

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;; APPLICANT: La Rosa Thomas J
;; APPLICANT: Kovalic David K
;; APPLICANT: Zhou Yihua
;; APPLICANT: Cao Yongwei
;; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
;; TITLE OF INVENTION: Placids and Uses Thereof for Plant Improvement
;; FILE REFERENCE: 38-21(53223)B
;; CURRENT APPLICATION NUMBER: US/10/424,599
;; CURRENT FILING DATE: 2003-04-28
;; NUMBER OF SEQ ID NOS: 285684
;; SEQ ID NO: 191823
;; LENGTH: 55
;; TYPE: PRT
;; ORGANISM: Glycine max
;; FEATURE:
;; OTHER INFORMATION: Clone ID: PAT_MRT3847_15236C.1.pcp
US-10-424-599-191823
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Query Match      78.3%; Score 36; DB 4; Length 55;
Best Local Similarity 87.5%; Pred. No. 34;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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```
Qy      1 LCTELQTT 8
        |||:||||
Db      13 LCMELQTT 20
```

```
RESULT 42
US-10-450-763-37868
; Sequence 37868, Application US/10450763
; Publication No. US20050196754A1
; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc
; TITLE OF INVENTION: NOVEL NUCLEIC ACIDS AND POLYPEPTIDES
; FILE REFERENCE: 790CIP3/US
; CURRENT APPLICATION NUMBER: US/10/450,763
; CURRENT FILING DATE: 2003-06-11
; PRIOR APPLICATION NUMBER: PCT/US01/08631
; PRIOR FILING DATE: 2001-03-30
; PRIOR APPLICATION NUMBER: 09/540,217
; PRIOR FILING DATE: 2000-03-31
; PRIOR APPLICATION NUMBER: 09/649,167
; PRIOR FILING DATE: 2000-08-23
; NUMBER OF SEQ ID NOS: 60736
; SOFTWARE: Custom
; SEQ ID NO: 37868
; LENGTH: 452
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (135)..(180)
; OTHER INFORMATION: kw TRANSCRIPTASE REVERSE II ORF2 domain identified by
; OTHER INFORMATION: EMATRIX, accession number DM01354N, p-value=1.000e-40, raw score
; OTHER INFORMATION: 13.17
US-10-450-763-37868
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Query Match      78.3%; Score 36; DB 5; Length 452;
Best Local Similarity 75.0%; Pred. No. 2,8e+02;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
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```
Qy      2 CTELQTTI 9
        |||:||||
Db      125 CSELQTTI 132
```

```
RESULT 43
US-10-408-765A-748
; Sequence 748, Application US/10408765A
; Publication No. US20040101874A1
; GENERAL INFORMATION:
; APPLICANT: Choeh, Soumitra S.
; APPLICANT: Fahy, Eoin D.
```

APPLICANT: Zhang, Bing
APPLICANT: Gibson, Bradford W.
APPLICANT: Taylor, Steven W.
APPLICANT: Glenn, Gary M.
APPLICANT: Marnock, Dale E.
TITLE OF INVENTION: TARGETS FOR THERAPEUTIC INTERVENTION
TITLE OF INVENTION: IDENTIFIED IN THE MITOCHONDRIAL PROTEOME
FILE REFERENCE: 660088.465
CURRENT APPLICATION NUMBER: US/10/408,765A
CURRENT FILING DATE: 2003-04-04
NUMBER OF SEQ ID NOS: 3077
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 748
LENGTH: 759
TYPE: PRT
ORGANISM: Homo sapiens
US-10-408-765A-748

Query Match 78.3%; Score 36; DB 4; Length 759;
Best Local Similarity 75.0%; Pred. No. 4.7e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LCTBLQTTI 9
DB 520 CTECQTV 527

RESULT 44
US-11-021-949-2
Sequence 2, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
TITLE OF INVENTION: AND METHODS OF THEIR USE
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 2
LENGTH: 25
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-2

Query Match 76.1%; Score 35; DB 6; Length 25;
Best Local Similarity 66.7%; Pred. No. 24;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTBLQTTI 9
DB 14 LCTBLQTTI 22

RESULT 45
US-11-021-949-57
Sequence 57, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
TITLE OF INVENTION: AND METHODS OF THEIR USE

FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 57
LENGTH: 25
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-57

Query Match 76.1%; Score 35; DB 6; Length 25;
Best Local Similarity 66.7%; Pred. No. 24;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTBLQTTI 9
DB 14 LCTBLQTTI 22

RESULT 46
US-10-751-845-152
Sequence 152, Application US/10751845
Publication No. US20050100928A1
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Urban, Robert G.
APPLICANT: Chicz, Roman M.
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
FILE REFERENCE: 08191-013001
CURRENT APPLICATION NUMBER: US/10/751,845
CURRENT FILING DATE: 2004-01-05
PRIOR APPLICATION NUMBER: US/09/664,225
PRIOR FILING DATE: 2000-08-18
PRIOR APPLICATION NUMBER: US 60/169,846
PRIOR FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: US 60/154,665
PRIOR FILING DATE: 1999-09-16
NUMBER OF SEQ ID NOS: 163
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 152
LENGTH: 42
TYPE: PRT
ORGANISM: Human Papilloma virus
US-10-751-845-152

Query Match 76.1%; Score 35; DB 5; Length 42;
Best Local Similarity 66.7%; Pred. No. 40;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTBLQTTI 9
DB 9 LCTBLQTTI 17

RESULT 47
US-10-437-963-175677
Sequence 175677, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated with
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B

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;; CURRENT APPLICATION NUMBER: US/10/437,963
;; CURRENT FILING DATE: 2003-05-14
;; NUMBER OF SEQ ID NOS: 204966
;; SEQ ID NO 175677
;; LENGTH: 82
;; TYPE: PR
;; ORGANISM: Oryza sativa
;; FEATURE:
;; OTHER INFORMATION: Clone ID: PAT_MRT4530_73499C.1.pep
US-10-437-963-175677

Query Match          76.1%; Score 35; DB 4; Length 82;
Best Local Similarity 75.0%; Pred. No. 78;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 CTELOTTI 9
   |||||
Db 38 CTELSL 45

RESULT 48
US-10-751-845-159
; Sequence 159, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Chicz, Robert G.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 159
; LENGTH: 119
; TYPE: PR
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Artificial fusion sequence
US-10-751-845-159

Query Match          76.1%; Score 35; DB 5; Length 119;
Best Local Similarity 66.7%; Pred. No. 1.1e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELOTTI 9
   |||||
Db 9 LCTELNTSL 17

RESULT 49
US-10-800-023-27
; Sequence 27, Application US/10800023
; Publication No. US20040256868A1
; GENERAL INFORMATION:
; APPLICANT: Steinman, Ralph
; APPLICANT: Nussenzweig, Michel
; APPLICANT: Hawiger, Daniel
; APPLICANT: Bonifaz, Laura
; TITLE OF INVENTION: Enhanced Antigen Delivery and Modulation
; TITLE OF INVENTION: Of the Immune Response Therefrom
; FILE REFERENCE: 600-1-081CONCIP1
; CURRENT APPLICATION NUMBER: US/10/800,023
; CURRENT FILING DATE: 2004-03-14
; PRIOR APPLICATION NUMBER: 09/925,284
; PRIOR FILING DATE: 2001-08-09
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;; PRIOR APPLICATION NUMBER: 09/586,704
;; PRIOR FILING DATE: 2000-06-05
;; PRIOR APPLICATION NUMBER: PCT/US96/01383
;; PRIOR FILING DATE: 1996-01-31
;; PRIOR APPLICATION NUMBER: 08/381,528
;; PRIOR FILING DATE: 1995-01-31
;; NUMBER OF SEQ ID NOS: 37
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 27
;; LENGTH: 158
;; TYPE: PR
;; ORGANISM: human papilloma virus E6 protein
US-10-800-023-27

Query Match          76.1%; Score 35; DB 5; Length 158;
Best Local Similarity 66.7%; Pred. No. 1.5e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELOTTI 9
   |||||
Db 17 LCTELNTSL 25

RESULT 50
US-11-021-949-28
; Sequence 28, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEITZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 28
; LENGTH: 158
; TYPE: PR
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-28

Query Match          76.1%; Score 35; DB 6; Length 158;
Best Local Similarity 66.7%; Pred. No. 1.5e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELOTTI 9
   |||||
Db 17 LCTELNTSL 25

Search completed: May 5, 2006, 08:55:36
Job time : 59 secs
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GenCore version 5.1.7
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OM protein - protein search, using SW model

Run on: May 5, 2006, 08:51:51 ; Search time 9 Seconds
(without alignments)
46.265 Million cell updates/sec

Title: US-08-170-344-4
Perfect score: 46
Sequence: 1 LCTELQTTI 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

Published Applications_AA_New:*
1: /SID5/ptodata/1/pubpaa/US08_NEW_PUB.pep1:*
2: /SID5/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
3: /SID5/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
4: /SID5/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
5: /SID5/ptodata/1/pubpaa/PTCT_NEW_PUB.pep:*
6: /SID5/ptodata/1/pubpaa/US09_NEW_PUB.pep:*
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12: /SID5/ptodata/1/pubpaa/US60_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	46	100.0	151	9 US-10-530-253-13	Sequence 13, Appl
2	46	100.0	158	11 US-11-206-138-3	Sequence 3, Appl
3	46	100.0	248	9 US-10-530-253-1	Sequence 1, Appl
4	46	100.0	248	9 US-10-530-253-3	Sequence 3, Appl
5	46	100.0	248	9 US-10-530-253-5	Sequence 5, Appl
6	46	100.0	248	9 US-10-530-253-7	Sequence 7, Appl
7	46	100.0	248	9 US-10-530-253-9	Sequence 9, Appl
8	46	100.0	248	9 US-10-530-253-11	Sequence 11, Appl
9	46	100.0	256	11 US-11-192-923A-2	Sequence 2, Appl
10	37	80.4	254	11 US-11-079-463-6325	Sequence 6325, Ap
11	35	76.1	158	9 US-10-530-253-15	Sequence 15, Appl
12	35	76.1	158	9 US-10-530-253-20	Sequence 20, Appl
13	35	73.9	587	11 US-11-188-296-296	Sequence 296, Ap
14	33	71.7	158	9 US-10-530-253-19	Sequence 19, Appl
15	33	71.7	420	11 US-11-087-099-6055	Sequence 6055, Ap
16	33	71.7	492	11 US-11-096-568A-14735	Sequence 14735, A
17	33	71.7	500	11 US-11-096-568A-14734	Sequence 14734, A
18	33	71.7	1310	11 US-11-096-568A-28552	Sequence 28552, A
19	33	71.7	1368	11 US-11-096-568A-28551	Sequence 28551, A
20	33	71.7	1374	11 US-11-096-568A-28550	Sequence 28550, A
21	31	67.4	43	9 US-10-467-657-7248	Sequence 7248, Ap

22	31	67.4	129	9 US-10-519-390-15	Sequence 15, Appl
23	31	67.4	128	11 US-11-176-830-207	Sequence 207, Ap
24	31	67.4	129	11 US-11-176-830-546	Sequence 546, Ap
25	31	67.4	129	11 US-11-176-830-547	Sequence 547, Ap
26	31	67.4	129	11 US-11-176-830-548	Sequence 548, Ap
27	31	67.4	129	11 US-11-176-830-549	Sequence 549, Ap
28	31	67.4	129	11 US-11-176-830-550	Sequence 550, Ap
29	31	67.4	129	11 US-11-176-830-551	Sequence 551, Ap
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31	31	67.4	129	11 US-11-176-830-553	Sequence 553, Ap
32	31	67.4	129	11 US-11-176-830-554	Sequence 554, Ap
33	31	67.4	129	11 US-11-176-830-555	Sequence 555, Ap
34	31	67.4	129	11 US-11-176-830-556	Sequence 556, Ap
35	31	67.4	129	11 US-11-176-830-557	Sequence 557, Ap
36	31	67.4	129	11 US-11-176-830-558	Sequence 558, Ap
37	31	67.4	129	11 US-11-176-830-559	Sequence 559, Ap
38	31	67.4	129	11 US-11-176-830-560	Sequence 560, Ap
39	31	67.4	129	11 US-11-176-830-561	Sequence 561, Ap
40	31	67.4	129	11 US-11-176-830-562	Sequence 562, Ap
41	31	67.4	129	11 US-11-176-830-563	Sequence 563, Ap
42	31	67.4	129	11 US-11-176-830-564	Sequence 564, Ap
43	31	67.4	129	11 US-11-176-830-565	Sequence 565, Ap
44	31	67.4	129	11 US-11-176-830-566	Sequence 566, Ap
45	31	67.4	129	11 US-11-176-830-567	Sequence 567, Ap
46	31	67.4	149	9 US-10-530-253-17	Sequence 17, Appl
47	31	67.4	150	9 US-10-469-561-25	Sequence 25, Appl
48	31	67.4	153	8 US-10-511-977-2476	Sequence 2476, Ap
49	31	67.4	153	11 US-11-174-398-8	Sequence 8, Appl
50	31	67.4	153	11 US-11-289-326-13	Sequence 13, Appl
51	31	67.4	255	11 US-11-026-396-5	Sequence 5, Appl
52	31	67.4	273	11 US-11-026-396-7	Sequence 7, Appl
53	31	67.4	293	11 US-11-026-396-4	Sequence 4, Appl
54	31	67.4	295	11 US-11-026-396-2	Sequence 2, Appl
55	31	67.4	412	11 US-11-087-099-5930	Sequence 5930, Ap
56	31	67.4	498	11 US-11-087-099-2684	Sequence 2684, Ap
57	31	67.4	514	11 US-11-087-099-2712	Sequence 2712, Ap
58	31	67.4	645	11 US-11-045-004-2516	Sequence 2516, Ap
59	31	67.4	929	9 US-10-467-657-6565	Sequence 5656, Ap
60	31	67.4	1320	11 US-11-098-686-1081	Sequence 10831, A
61	31	67.4	4913	9 US-10-453-372-1142	Sequence 1142, Ap
62	31	67.4	9	US-10-453-372-1132	Sequence 1132, Ap
63	30	65.2	22	11 US-11-207-078-280	Sequence 280, Ap
64	30	65.2	34	11 US-11-207-078-256	Sequence 256, Ap
65	30	65.2	42	11 US-11-207-078-96	Sequence 96, Appl
66	30	65.2	47	11 US-11-207-078-42	Sequence 42, Appl
67	30	65.2	126	11 US-11-096-568A-3394	Sequence 3394, Ap
68	30	65.2	146	11 US-11-096-568A-14842	Sequence 14842, A
69	30	65.2	157	11 US-11-207-078-126	Sequence 126, Ap
70	30	65.2	160	9 US-10-846-172A-10	Sequence 10, Appl
71	30	65.2	196	11 US-11-120-925-120	Sequence 120, Ap
72	30	65.2	202	11 US-11-098-662-8	Sequence 8, Appl
73	30	65.2	202	11 US-11-098-662-10	Sequence 10, Appl
74	30	65.2	202	11 US-11-193-955-8	Sequence 8, Appl
75	30	65.2	202	11 US-11-193-955-10	Sequence 10, Appl
76	30	65.2	202	11 US-11-266-626-9	Sequence 9, Appl
77	30	65.2	202	11 US-11-266-626-11	Sequence 11, Appl
78	30	65.2	215	11 US-11-045-004-1613	Sequence 1613, Ap
79	30	65.2	231	11 US-11-096-568A-3393	Sequence 3393, Ap
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81	30	65.2	631	11 US-11-188-298-16078	Sequence 16078, A
82	30	65.2	884	11 US-11-207-078-222	Sequence 222, Ap
83	30	65.2	1194	11 US-11-098-666-10270	Sequence 10270, A
84	30	65.2	1410	9 US-10-821-23A-1050	Sequence 1050, Ap
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87	29	63.0	174	11 US-11-087-099-1876	Sequence 1876, Ap
88	29	63.0	175	11 US-11-087-099-9020	Sequence 9020, Ap
89	29	63.0	175	11 US-11-096-568A-4212	Sequence 4212, Ap
90	29	63.0	197	11 US-11-172-740-2085	Sequence 2085, Ap
91	29	63.0	214	11 US-11-096-568A-4211	Sequence 4211, Ap
92	29	63.0	222	11 US-11-087-099-6133	Sequence 6133, Ap
93	29	63.0	284	11 US-11-212-443-58	Sequence 58, Appl
94	29	63.0	284	11 US-11-087-099-10570	Sequence 10570, A

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96	29	63.0	307	9	US-10-055-877-232	Sequence 232, App	169	28	60.9	2644	9	US-10-770-726-45	Sequence 45, Appl
97	29	63.0	360	11	US-11-212-443-178	Sequence 178, App	170	27	59.8	300	11	US-11-188-298-10963	Sequence 10963, A
98	29	63.0	400	11	US-11-188-298-22075	Sequence 22075, A	171	27	58.7	19	11	US-11-004-399-1357	Sequence 1357, Ap
99	29	63.0	405	11	US-11-188-298-15166	Sequence 15166, A	172	27	58.7	35	9	US-10-821-234-1704	Sequence 1704, Ap
100	29	63.0	452	11	US-11-087-099-4768	Sequence 4768, A	173	27	58.7	65	11	US-11-096-5688-27328	Sequence 27328, A
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102	29	63.0	484	11	US-11-087-099-5622	Sequence 6203, Ap	175	27	58.7	84	11	US-11-194-246-295	Sequence 295, App
103	29	63.0	484	11	US-11-096-5688-6203	Sequence 8260, Ap	176	27	58.7	137	11	US-11-096-5688-5527	Sequence 5227, Ap
104	29	63.0	489	11	US-11-079-463-8260	Sequence 6202, Ap	177	27	58.7	140	11	US-11-072-512-2719	Sequence 2719, Ap
105	29	63.0	499	11	US-11-096-5688-6202	Sequence 6201, Ap	178	27	58.7	157	9	US-10-980-388-78	Sequence 78, Appl
106	29	63.0	520	11	US-11-087-099-5096	Sequence 5096, Ap	179	27	58.7	158	9	US-10-530-253-26	Sequence 26, Appl
107	29	63.0	1532	11	US-11-212-443-62	Sequence 62, Appl	180	27	58.7	168	9	US-10-454-437-196	Sequence 196, App
108	29	63.0	1542	9	US-10-453-372-258	Sequence 258, App	181	27	58.7	168	11	US-11-055-822-360	Sequence 360, App
109	29	63.0	1542	9	US-10-453-372-266	Sequence 266, App	182	27	58.7	189	11	US-11-096-5688-6726	Sequence 6726, Ap
110	29	63.0	1542	9	US-10-453-372-280	Sequence 280, App	183	27	58.7	201	11	US-11-096-5688-2933	Sequence 2939, Ap
111	29	63.0	1574	11	US-11-212-443-179	Sequence 179, App	184	27	58.7	211	11	US-11-087-099-10457	Sequence 10457, A
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113	29	63.0	1796	11	US-11-188-298-9498	Sequence 9498, Ap	186	27	58.7	212	11	US-11-188-298-1528	Sequence 1528, Ap
114	29	63.0	1978	11	US-11-212-443-60	Sequence 60, Appl	187	27	58.7	214	11	US-11-096-5688-6725	Sequence 6725, Ap
115	29	63.0	1981	11	US-11-045-208-38	Sequence 38, Appl	188	27	58.7	231	9	US-10-915-002-347	Sequence 347, App
116	29	63.0	2015	11	US-11-052-5544-374	Sequence 374, Appl	189	27	58.7	231	9	US-10-915-002-351	Sequence 351, App
117	28	60.9	12	11	US-11-103-356A-12	Sequence 12, Appl	190	27	58.7	231	11	US-11-096-5688-2838	Sequence 2938, Ap
118	28	60.9	115	11	US-11-096-5688-31408	Sequence 31408, A	191	27	58.7	238	11	US-11-098-686-10627	Sequence 10627, A
119	28	60.9	118	11	US-11-072-512-3683	Sequence 3683, Ap	192	27	58.7	267	11	US-11-098-686-10320	Sequence 10320, A
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122	28	60.9	172	11	US-11-188-298-17891	Sequence 17891, A	195	27	58.7	276	9	US-10-251-567-1	Sequence 1, Appl1
123	28	60.9	198	9	US-10-467-657-4664	Sequence 4664, Ap	196	27	58.7	278	11	US-11-233-798-12	Sequence 12, Appl
124	28	60.9	252	11	US-11-229-769-214	Sequence 214, App	197	27	58.7	287	11	US-11-096-5688-2337	Sequence 2937, Ap
125	28	60.9	267	11	US-11-140-024-5	Sequence 5, Appl1	198	27	58.7	288	9	US-10-525-907-44	Sequence 44, Appl1
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127	28	60.9	277	8	US-10-511-937-2455	Sequence 2455, Ap	200	27	58.7	297	11	US-11-079-463-8280	Sequence 8280, Ap
128	28	60.9	277	11	US-11-132-285-3	Sequence 3, Appl1	201	27	58.7	314	11	US-11-188-298-641	Sequence 6471, Ap
129	28	60.9	277	11	US-11-182-946-12	Sequence 12, Appl1	202	27	58.7	317	11	US-11-129-143-85	Sequence 85, Appl
130	28	60.9	308	11	US-11-098-686-11286	Sequence 11286, A	203	27	58.7	317	11	US-11-096-5688-7806	Sequence 7806, Ap
131	28	60.9	334	11	US-11-098-686-10496	Sequence 10496, A	204	27	58.7	317	11	US-11-188-298-2979	Sequence 2979, Ap
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133	28	60.9	366	11	US-11-087-099-4201	Sequence 4201, Ap	206	27	58.7	326	11	US-11-000-463-382	Sequence 382, App
134	28	60.9	366	11	US-11-087-099-7455	Sequence 7455, Ap	207	27	58.7	326	11	US-11-000-463-854	Sequence 854, App
135	28	60.9	366	11	US-11-087-099-9353	Sequence 9353, Ap	208	27	58.7	326	11	US-11-096-5688-1102	Sequence 1202, Ap
136	28	60.9	368	11	US-11-087-099-936	Sequence 936, App	209	27	58.7	328	11	US-11-096-5688-3897	Sequence 3897, Ap
137	28	60.9	368	11	US-11-087-099-4522	Sequence 4522, Ap	210	27	58.7	334	11	US-11-096-5688-3896	Sequence 3896, Ap
138	28	60.9	368	11	US-11-087-099-9713	Sequence 9713, Ap	211	27	58.7	334	11	US-11-172-740-45	Sequence 45, Appl1
139	28	60.9	387	11	US-11-096-5688-25143	Sequence 25143, A	212	27	58.7	335	11	US-11-188-298-16004	Sequence 16004, A
140	28	60.9	389	11	US-11-096-5688-25142	Sequence 25142, A	213	27	58.7	335	11	US-11-288-493-58	Sequence 58, Appl1
141	28	60.9	413	8	US-10-511-937-2428	Sequence 2428, A	214	27	58.7	337	11	US-11-188-298-6332	Sequence 6382, Appl
142	28	60.9	413	9	US-10-880-764-3	Sequence 3, Appl1	215	27	58.7	337	11	US-11-188-298-13359	Sequence 13359, A
143	28	60.9	432	11	US-11-096-5688-25141	Sequence 25141, A	216	27	58.7	343	11	US-11-096-5688-6730	Sequence 6730, Ap
144	28	60.9	446	11	US-11-096-5688-19198	Sequence 19198, A	217	27	58.7	344	11	US-11-079-463-6695	Sequence 6695, Ap
145	28	60.9	451	11	US-11-096-5688-14775	Sequence 14775, A	218	27	58.7	346	11	US-11-096-5688-1201	Sequence 1201, Ap
146	28	60.9	480	11	US-11-096-5688-14775	Sequence 15197, A	219	27	58.7	352	11	US-11-096-5688-6729	Sequence 6729, Ap
147	28	60.9	601	11	US-11-024-959-295	Sequence 295, App	220	27	58.7	365	11	US-11-229-769-332	Sequence 322, App
148	28	60.9	719	9	US-10-511-538-247	Sequence 247, App	221	27	58.7	371	11	US-11-188-298-16216	Sequence 16216, A
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151	28	60.9	1098	11	US-11-072-512-2475	Sequence 2475, Ap	224	27	58.7	388	11	US-11-188-298-21443	Sequence 21843, A
152	28	60.9	1220	9	US-10-501-035-260	Sequence 260, App	225	27	58.7	401	9	US-10-878-556A-179	Sequence 179, App
153	28	60.9	1828	11	US-11-126-022-23	Sequence 22, Appl	226	27	58.7	401	11	US-11-188-298-9920	Sequence 9920, Ap
154	28	60.9	1852	11	US-11-126-022-1	Sequence 21, Appl	227	27	58.7	402	11	US-11-188-298-7740	Sequence 7740, Ap
155	28	60.9	1863	11	US-11-126-022-2	Sequence 2, Appl1	228	27	58.7	421	8	US-10-505-928-110	Sequence 410, App
156	28	60.9	1863	11	US-11-126-022-9	Sequence 9, Appl1	229	27	58.7	421	9	US-10-878-556A-173	Sequence 173, App
157	28	60.9	1863	11	US-11-126-022-10	Sequence 10, Appl	230	27	58.7	437	11	US-11-087-099-5721	Sequence 5721, Ap
158	28	60.9	1863	11	US-11-126-022-11	Sequence 11, Appl	231	27	58.7	437	11	US-11-188-298-5230	Sequence 5230, Ap
159	28	60.9	1863	11	US-11-126-022-12	Sequence 12, Appl	232	27	58.7	447	11	US-11-098-686-11121	Sequence 1121, A
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161	28	60.9	1863	11	US-11-126-022-14	Sequence 14, Appl	234	27	58.7	447	11	US-11-188-298-7480	Sequence 7480, Ap
162	28	60.9	1863	11	US-11-126-022-15	Sequence 15, Appl	235	27	58.7	447	11	US-11-188-298-18043	Sequence 18043, A
163	28	60.9	1863	11	US-11-126-022-16	Sequence 16, Appl	236	27	58.7	449	9	US-10-821-234-1089	Sequence 1089, Ap
164	28	60.9	1863	11	US-11-126-022-17	Sequence 17, Appl	237	27	58.7	451	11	US-11-188-298-5939	Sequence 5939, Ap
165	28	60.9	1863	11	US-11-126-022-18	Sequence 18, Appl	238	27	58.7	457	11	US-11-087-099-10361	Sequence 10361, A
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242	27	58.7	542	11	US-11-229-769-323	Sequence 323, App	315	26	56.5	284	11	US-11-102-240-62	Sequence 62, Appl
243	27	58.7	545	11	US-11-188-298-16184	Sequence 16184, A	316	26	56.5	284	11	US-11-103-195-62	Sequence 62, Appl
244	27	58.7	545	11	US-11-096-568A-1994	Sequence 1994, Ap	317	26	56.5	298	11	US-11-096-568A-1942	Sequence 1942, Ap
245	27	58.7	565	11	US-11-096-568A-27236	Sequence 27236, A	318	26	56.5	303	11	US-11-096-568A-1941	Sequence 89, Appl
246	27	58.7	570	11	US-11-113-424-69	Sequence 69, Appl	319	26	56.5	306	11	US-11-019-711-89	Sequence 90, Appl
247	27	58.7	570	11	US-11-113-424-71	Sequence 71, Appl	320	26	56.5	306	11	US-11-019-711-90	Sequence 90, Appl
248	27	58.7	620	11	US-11-113-424-70	Sequence 70, Appl	321	26	56.5	307	10	US-11-283-522-33	Sequence 87, Appl
249	27	58.7	624	11	US-11-096-568A-1993	Sequence 1993, Ap	322	26	56.5	307	11	US-11-019-711-88	Sequence 87, Appl
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251	27	58.7	625	11	US-11-096-568A-1992	Sequence 1992, Ap	324	26	56.5	307	11	US-11-265-966-16	Sequence 16, Appl
252	27	58.7	625	11	US-11-024-959-502	Sequence 502, App	325	26	56.5	309	11	US-11-087-099-11588	Sequence 10246, A
253	27	58.7	641	11	US-11-024-959-502	Sequence 4572, Ap	326	26	56.5	309	11	US-11-087-099-11588	Sequence 11588, A
254	27	58.7	691	11	US-11-087-099-4572	Sequence 26, Appl	327	26	56.5	310	11	US-11-096-568A-6851	Sequence 6851, Ap
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256	27	58.7	734	9	US-10-501-035-347	Sequence 26, Appl	329	26	56.5	320	11	US-11-087-099-11634	Sequence 11834, A
257	27	58.7	747	11	US-11-113-424-26	Sequence 774, App	330	26	56.5	326	11	US-11-096-568A-15425	Sequence 15425, A
258	27	58.7	747	11	US-11-264-096-774	Sequence 648, App	331	26	56.5	327	11	US-11-072-512-2508	Sequence 2839, Ap
259	27	58.7	762	9	US-10-506-454-648	Sequence 316, App	332	26	56.5	330	11	US-11-072-512-2839	Sequence 2839, Ap
260	27	58.7	821	8	US-10-505-928-316	Sequence 316, App	333	26	56.5	330	11	US-11-087-099-5040	Sequence 5040, Ap
261	27	58.7	821	11	US-11-087-227-90	Sequence 90, Appl	334	26	56.5	331	11	US-11-188-298-13124	Sequence 13124, A
262	27	58.7	821	11	US-11-233-510-16	Sequence 16, Appl	335	26	56.5	336	11	US-11-096-568A-13760	Sequence 12760, A
263	27	58.7	1014	11	US-11-188-298-10090	Sequence 10090, A	336	26	56.5	339	11	US-11-096-568A-15425	Sequence 15425, A
264	27	58.7	1095	9	US-10-793-628-3154	Sequence 3154, Ap	337	26	56.5	339	11	US-11-172-740-326	Sequence 326, App
265	27	58.7	1275	9	US-10-821-234-1598	Sequence 1598, Ap	338	26	56.5	339	11	US-11-096-568A-6850	Sequence 6850, Ap
266	27	58.7	1663	9	US-10-055-877-148	Sequence 148, App	339	26	56.5	344	11	US-11-087-099-720	Sequence 720, App
267	27	58.7	1778	11	US-11-087-099-12160	Sequence 12160, A	340	26	56.5	346	11	US-11-087-099-6518	Sequence 6518, Ap
268	27	58.7	1778	11	US-11-045-004-19	Sequence 19, Appl	341	26	56.5	346	11	US-11-087-099-1693	Sequence 33378, A
269	27	58.7	1985	9	US-10-501-035-218	Sequence 218, App	342	26	56.5	348	11	US-11-096-568A-13378	Sequence 1693, Ap
270	26.5	57.6	146	11	US-11-096-568A-29968	Sequence 29968, A	343	26	56.5	349	11	US-11-087-099-1693	Sequence 1693, Ap
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272	26.5	57.6	170	11	US-11-096-568A-29966	Sequence 29966, A	345	26	56.5	350	11	US-11-188-298-21850	Sequence 21850, A
273	26.5	57.6	261	11	US-11-079-463-8265	Sequence 8265, Ap	346	26	56.5	352	11	US-11-096-568A-22483	Sequence 22483, A
274	26	56.5	19	11	US-11-004-399-991	Sequence 991, App	347	26	56.5	353	11	US-11-096-568A-10134	Sequence 10134, A
275	26	56.5	49	9	US-10-895-064-2872	Sequence 2872, Ap	348	26	56.5	355	11	US-10-506-454-16	Sequence 16, Appl
276	26	56.5	49	11	US-11-123-741-8872	Sequence 2872, Ap	349	26	56.5	356	9	US-10-288-733-2	Sequence 2, Appl
277	26	56.5	55	11	US-11-096-568A-7906	Sequence 7906, Ap	350	26	56.5	360	9	US-11-188-298-10162	Sequence 10162, A
278	26	56.5	90	11	US-11-264-096-1984	Sequence 12, Appl	351	26	56.5	368	11	US-11-096-568A-31554	Sequence 31554, A
279	26	56.5	108	11	US-11-188-298-5279	Sequence 1984, Ap	352	26	56.5	373	11	US-11-055-822-458	Sequence 458, App
280	26	56.5	116	11	US-11-176-830-544	Sequence 5279, Ap	353	26	56.5	375	11	US-11-087-099-1201	Sequence 1201, Ap
281	26	56.5	129	11	US-11-176-830-544	Sequence 544, App	354	26	56.5	375	11	US-11-087-099-1201	Sequence 7023, Ap
282	26	56.5	129	11	US-11-176-830-545	Sequence 544, App	355	26	56.5	375	11	US-11-188-298-12517	Sequence 12517, A
283	26	56.5	130	11	US-11-045-004-977	Sequence 977, App	356	26	56.5	375	11	US-11-087-099-7023	Sequence 31553, A
284	26	56.5	142	11	US-11-079-463-8099	Sequence 8099, Ap	357	26	56.5	378	11	US-11-096-568A-31553	Sequence 6334, A
285	26	56.5	148	8	US-10-505-928-237	Sequence 237, App	358	26	56.5	379	11	US-11-188-298-6334	Sequence 6334, A
286	26	56.5	149	9	US-10-530-253-24	Sequence 24, Appl	359	26	56.5	381	11	US-11-188-298-15196	Sequence 15196, A
287	26	56.5	164	11	US-11-045-004-423	Sequence 423, App	360	26	56.5	384	9	US-10-525-674-44	Sequence 44, Appl
288	26	56.5	172	9	US-10-793-626-1266	Sequence 1266, Ap	361	26	56.5	385	11	US-11-045-004-1508	Sequence 1508, Ap
289	26	56.5	172	11	US-11-098-686-10527	Sequence 10527, A	362	26	56.5	389	11	US-11-096-568A-19737	Sequence 19737, A
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291	26	56.5	177	11	US-11-072-512-3347	Sequence 3347, Ap	364	26	56.5	396	11	US-11-096-568A-19736	Sequence 19736, A
292	26	56.5	181	11	US-11-072-512-3327	Sequence 3327, Ap	365	26	56.5	396	11	US-11-188-298-10645	Sequence 10645, A
293	26	56.5	181	11	US-11-264-096-910	Sequence 910, App	366	26	56.5	396	11	US-11-188-298-17724	Sequence 17724, A
294	26	56.5	212	11	US-11-098-686-11253	Sequence 11253, A	367	26	56.5	398	11	US-11-072-512-3388	Sequence 3388, Ap
295	26	56.5	214	11	US-11-096-568A-4858	Sequence 4858, Ap	368	26	56.5	400	11	US-11-188-298-5081	Sequence 5081, Ap
296	26	56.5	215	11	US-11-188-298-2653	Sequence 2653, Ap	369	26	56.5	401	11	US-11-096-568A-33377	Sequence 33377, A
297	26	56.5	215	11	US-11-1202-516-6	Sequence 6, Appl	370	26	56.5	405	11	US-11-188-298-9774	Sequence 9774, Ap
298	26	56.5	215	11	US-11-238-936-12	Sequence 12, Appl	371	26	56.5	405	11	US-11-188-298-9774	Sequence 11528, A
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308	26	56.5	271	11	US-11-019-711-26	Sequence 26, Appl	381	26	56.5	443	9	US-10-131-826A-318	Sequence 318, App
309	26	56.5	282	11	US-11-096-568A-4857	Sequence 4857, Ap	382	26	56.5	443	9	US-10-973-115B-118	Sequence 318, App
310	26	56.5	284	9	US-10-063-703-62	Sequence 62, Appl	383	26	56.5	443	9	US-10-137-872A-318	Sequence 318, App
311	26	56.5	284	9	US-10-194-487-236	Sequence 236, App	384	26	56.5	443	9	US-10-152-370-318	Sequence 318, App
312	26	56.5	284	9	US-10-195-883-236	Sequence 236, App	385	26	56.5	443	9	US-10-152-370-318	Sequence 318, App
313	26	56.5	284	9	US-10-195-888-236	Sequence 236, App	386	26	56.5	443	11	US-11-065-695-16	Sequence 16, Appl

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389	26	56.5	444	9	US-10-513-639-22	Sequence 22, Appl	462	26	56.5	941	11	US-11-124-367A-315	Sequence 315, App
390	26	56.5	444	9	US-10-513-639-23	Sequence 23, Appl	463	26	56.5	1075	11	US-11-100-640-12	Sequence 12, Appl
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392	26	56.5	451	11	US-11-188-298-951	Sequence 951, App	465	26	56.5	1083	11	US-11-113-751-40	Sequence 40, Appl
393	26	56.5	462	11	US-11-124-367A-497	Sequence 497, App	466	26	56.5	1110	11	US-11-113-751-4	Sequence 4, Appl1
394	26	56.5	464	9	US-10-689-742-164	Sequence 164, App	467	26	56.5	1115	11	US-11-113-751-14	Sequence 14, Appl
395	26	56.5	464	11	US-11-188-298-5875	Sequence 5875, App	468	26	56.5	1115	11	US-11-113-751-46	Sequence 46, Appl
396	26	56.5	464	11	US-11-188-298-12067	Sequence 12067, A	469	26	56.5	1116	11	US-11-113-751-32	Sequence 32, Appl
397	26	56.5	468	11	US-11-188-298-2846	Sequence 2846, Ap	470	26	56.5	1116	11	US-11-113-751-34	Sequence 34, Appl
398	26	56.5	468	11	US-11-188-298-9547	Sequence 9547, Ap	471	26	56.5	1118	11	US-11-113-751-42	Sequence 42, Appl
399	26	56.5	471	11	US-11-096-568A-15988	Sequence 15988, A	472	26	56.5	1121	11	US-11-113-751-19	Sequence 19, Appl
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401	26	56.5	472	11	US-11-188-298-1709	Sequence 7009, Ap	474	26	56.5	1151	11	US-11-113-751-36	Sequence 36, Appl
402	26	56.5	472	11	US-11-188-298-7690	Sequence 7690, Ap	475	26	56.5	1153	11	US-11-113-751-44	Sequence 44, Appl
403	26	56.5	472	11	US-11-188-298-9346	Sequence 9346, Ap	476	26	56.5	1165	11	US-11-188-298-17834	Sequence 44, Appl
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405	26	56.5	474	11	US-11-096-568A-26453	Sequence 26453, A	478	26	56.5	1255	11	US-11-022-562-235	Sequence 235, App
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407	26	56.5	484	11	US-11-096-568A-15987	Sequence 15987, A	480	26	56.5	1255	11	US-11-052-554A-266	Sequence 266, App
408	26	56.5	486	11	US-11-072-512-3837	Sequence 3837, Ap	481	26	56.5	1255	11	US-11-052-554A-267	Sequence 267, App
409	26	56.5	489	11	US-11-072-512-2922	Sequence 2922, Ap	482	26	56.5	1255	11	US-11-052-554A-268	Sequence 268, App
410	26	56.5	495	11	US-11-198-886-15	Sequence 15, Appl	483	26	56.5	1255	11	US-11-052-554A-269	Sequence 269, App
411	26	56.5	498	11	US-11-131-479-2	Sequence 2, Appl1	484	26	56.5	1255	11	US-11-052-554A-270	Sequence 270, App
412	26	56.5	498	11	US-11-131-479-76	Sequence 76, Appl	485	26	56.5	1255	11	US-11-052-554A-271	Sequence 271, App
413	26	56.5	507	11	US-11-188-298-1536	Sequence 1536, Ap	486	26	56.5	1255	11	US-11-052-554A-272	Sequence 272, App
414	26	56.5	515	11	US-11-188-298-21436	Sequence 21436, A	487	26	56.5	1255	11	US-11-052-554A-273	Sequence 273, App
415	26	56.5	522	11	US-11-131-479-7	Sequence 7, Appl1	488	26	56.5	1255	11	US-11-052-554A-274	Sequence 274, App
416	26	56.5	522	11	US-11-131-479-9	Sequence 9, Appl1	489	26	56.5	1255	11	US-11-052-554A-275	Sequence 275, App
417	26	56.5	524	11	US-11-118-809-4	Sequence 4, Appl1	490	26	56.5	1255	11	US-11-004-399-4026	Sequence 4026, Ap
418	26	56.5	528	11	US-11-118-809-2	Sequence 2, Appl1	491	26	56.5	1257	11	US-11-004-399-1737	Sequence 1737, Ap
419	26	56.5	533	11	US-11-096-568A-20042	Sequence 20042, A	492	26	56.5	1279	9	US-10-957-880-3	Sequence 3, Appl1
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421	26	56.5	546	11	US-11-124-367A-499	Sequence 499, App	494	26	56.5	1441	11	US-11-096-568A-34298	Sequence 34298, A
422	26	56.5	549	11	US-11-188-298-8346	Sequence 8346, Ap	495	26	56.5	1450	11	US-11-096-568A-34297	Sequence 34297, A
423	26	56.5	549	11	US-11-188-298-15264	Sequence 15264, A	496	26	56.5	1488	11	US-11-096-568A-34296	Sequence 34296, A
424	26	56.5	553	11	US-11-103-857-61	Sequence 61, Appl	497	26	56.5	2244	11	US-11-131-263-9	Sequence 9, Appl1
425	26	56.5	554	11	US-11-055-822-998	Sequence 98, App	498	26	56.5	2244	11	US-11-131-263-18	Sequence 18, Appl
426	26	56.5	556	9	US-10-506-454-828	Sequence 828, App	499	26	56.5	2244	11	US-11-131-263-30	Sequence 30, Appl
427	26	56.5	590	11	US-11-087-099-8075	Sequence 8075, Ap	500	26	56.5	2244	11	US-11-131-263-41	Sequence 41, Appl
428	26	56.5	591	11	US-11-087-099-5735	Sequence 5735, Ap	501	26	56.5	3803	9	US-10-995-561-773	Sequence 713, Appl
429	26	56.5	611	11	US-11-188-298-5827	Sequence 5827, Ap	502	26	56.5	3960	9	US-10-995-561-771	Sequence 711, App
430	26	56.5	613	11	US-11-087-099-9717	Sequence 9717, Ap	503	26	56.5	5335	9	US-10-995-561-777	Sequence 777, App
431	26	56.5	640	11	US-11-087-099-4980	Sequence 4980, Ap	504	26	56.5	5406	9	US-10-995-561-774	Sequence 774, App
432	26	56.5	657	9	US-10-957-880-4	Sequence 4, Appl1	505	26	56.5	5415	9	US-10-995-561-779	Sequence 779, App
433	26	56.5	664	9	US-10-542-178-3	Sequence 3, Appl1	506	26	56.5	5465	9	US-10-995-561-775	Sequence 775, App
434	26	56.5	669	9	US-10-997-201A-30	Sequence 289, App	507	26	56.5	5935	9	US-10-995-561-776	Sequence 776, App
435	26	56.5	676	11	US-11-004-399-289	Sequence 3207, Ap	508	25	55.4	136	11	US-11-156-084-324	Sequence 324, App
436	26	56.5	676	11	US-11-004-399-3207	Sequence 3207, Ap	509	25	55.4	244	11	US-11-156-084-218	Sequence 218, App
437	26	56.5	694	11	US-11-188-298-19841	Sequence 19841, A	510	25	54.3	8	11	US-11-045-024-9858	Sequence 9858, Ap
438	26	56.5	701	11	US-11-188-298-19841	Sequence 1, Appl1	511	25	54.3	8	11	US-11-045-024-9858	Sequence 9858, Ap
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443	26	56.5	754	9	US-10-455-772-294	Sequence 294, App	516	25	54.3	10	11	US-11-045-024-9664	Sequence 9684, Ap
444	26	56.5	754	9	US-10-455-772-298	Sequence 298, App	517	25	54.3	10	11	US-11-045-024-9934	Sequence 9934, Ap
445	26	56.5	755	9	US-10-455-772-290	Sequence 290, App	518	25	54.3	10	11	US-11-045-024-11685	Sequence 11685, A
446	26	56.5	755	9	US-10-455-772-302	Sequence 302, App	519	25	54.3	10	11	US-11-045-024-11866	Sequence 11866, A
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448	26	56.5	756	9	US-10-455-772-296	Sequence 296, App	521	25	54.3	11	11	US-11-045-024-1371	Sequence 3371, Ap
449	26	56.5	777	9	US-10-455-772-538	Sequence 538, App	522	25	54.3	11	11	US-11-045-024-9700	Sequence 9700, Ap
450	26	56.5	778	9	US-10-455-772-292	Sequence 292, App	523	25	54.3	11	11	US-11-045-024-9914	Sequence 9914, Ap
451	26	56.5	778	11	US-11-264-096-912	Sequence 912, App	524	25	54.3	11	11	US-11-045-024-11693	Sequence 11693, A
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453	26	56.5	830	9	US-10-330-773-404	Sequence 406, App	526	25	54.3	15	11	US-11-045-024-13433	Sequence 13433, A
454	26	56.5	830	9	US-10-330-773-406	Sequence 406, App	527	25	54.3	15	11	US-11-121-361-18	Sequence 18, Appl
455	26	56.5	834	9	US-10-501-035-280	Sequence 280, App	528	25	54.3	34	11	US-11-096-568A-25411	Sequence 25411, A
456	26	56.5	851	9	US-10-330-773-397	Sequence 397, App	529	25	54.3	56	11	US-11-188-298-2953	Sequence 2953, Ap
457	26	56.5	868	9	US-10-821-234-1082	Sequence 1082, App	530	25	54.3	67	11	US-11-096-568A-2425	Sequence 2425, Ap
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459	26	56.5	885	11	US-11-096-568A-30498	Sequence 30499, A	532	25	54.3	79	11	US-11-264-096-406	Sequence 406, App

533	25	54.3	79	11	US-11-264-096-407	Sequence 407, App	606	25	54.3	268	11	US-11-072-512-2981	Sequence 2981, App
534	25	54.3	80	11	US-11-096-568A-2424	Sequence 2424, App	607	25	54.3	268	11	US-11-096-568A-4454	Sequence 4454, App
535	25	54.3	82	11	US-11-096-568A-1519	Sequence 1519, App	608	25	54.3	270	11	US-11-098-686-11113	Sequence 11113, A
536	25	54.3	91	11	US-11-082-389-352	Sequence 352, App	609	25	54.3	276	11	US-11-087-099-10298	Sequence 10298, A
537	25	54.3	102	11	US-11-264-096-1133	Sequence 1133, App	610	25	54.3	283	11	US-11-096-568A-5964	Sequence 5964, App
538	25	54.3	102	11	US-11-264-096-1185	Sequence 1185, App	611	25	54.3	283	11	US-11-079-463-8546	Sequence 8546, App
539	25	54.3	108	11	US-11-096-568A-3660	Sequence 3660, App	612	25	54.3	284	11	US-11-188-298-579	Sequence 579, App
540	25	54.3	113	11	US-11-072-512-2416	Sequence 2416, App	613	25	54.3	286	11	US-11-188-298-13264	Sequence 13264, A
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542	25	54.3	117	11	US-11-096-568A-3659	Sequence 3659, App	615	25	54.3	289	11	US-11-096-568A-30765	Sequence 30765, A
543	25	54.3	123	11	US-11-098-686-11187	Sequence 11187, A	616	25	54.3	289	11	US-11-188-298-8105	Sequence 8105, App
544	25	54.3	124	11	US-11-194-246-397	Sequence 397, App	617	25	54.3	291	11	US-11-098-686-11384	Sequence 11384, A
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546	25	54.3	127	11	US-11-197-380-2	Sequence 2, App1	619	25	54.3	295	9	US-10-784-004-408	Sequence 408, App
547	25	54.3	127	11	US-11-197-380-4	Sequence 4, App1	620	25	54.3	295	10	US-10-784-004-428	Sequence 728, App
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550	25	54.3	131	11	US-11-188-298-5337	Sequence 5337, App	623	25	54.3	299	11	US-11-058-924-10	Sequence 10, App1
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553	25	54.3	139	9	US-10-467-657-7880	Sequence 7880, App	626	25	54.3	303	11	US-11-172-740-1720	Sequence 1720, App
554	25	54.3	140	9	US-10-967-093-6	Sequence 4, App1	627	25	54.3	303	11	US-11-045-004-1069	Sequence 1069, App
555	25	54.3	140	9	US-10-967-093-6	Sequence 6, App1	628	25	54.3	304	11	US-11-098-686-10872	Sequence 10872, A
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557	25	54.3	144	11	US-11-096-568A-21463	Sequence 21463, A	630	25	54.3	307	9	US-10-793-626-2122	Sequence 2122, App
558	25	54.3	149	9	US-10-530-253-18	Sequence 18, App1	631	25	54.3	311	9	US-10-467-657-5644	Sequence 5644, App
559	25	54.3	155	11	US-11-096-568A-3658	Sequence 3658, App	632	25	54.3	316	11	US-11-058-924-2	Sequence 2, App1
560	25	54.3	157	11	US-11-188-298-1481	Sequence 1481, App	633	25	54.3	316	11	US-11-096-568A-25115	Sequence 25115, A
561	25	54.3	161	11	US-11-096-568A-31524	Sequence 31524, A	634	25	54.3	320	11	US-11-096-568A-21383	Sequence 21383, A
562	25	54.3	170	11	US-11-079-463-9969	Sequence 9969, App	635	25	54.3	326	11	US-11-229-371-75	Sequence 75, App1
563	25	54.3	171	11	US-11-098-686-11167	Sequence 11167, A	636	25	54.3	326	11	US-11-072-512-3428	Sequence 3428, App
564	25	54.3	172	11	US-11-177-010-2	Sequence 2, App1	637	25	54.3	326	11	US-11-228-823-75	Sequence 75, App1
565	25	54.3	172	11	US-11-177-010-4	Sequence 4, App1	638	25	54.3	326	11	US-11-228-875-75	Sequence 75, App1
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574	25	54.3	182	9	US-10-453-372-1104	Sequence 1104, App	647	25	54.3	335	11	US-11-188-298-11875	Sequence 11875, A
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587	25	54.3	227	9	US-10-527-771-12	Sequence 12, App1	660	25	54.3	369	11	US-11-087-099-7687	Sequence 7687, App
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597	25	54.3	247	11	US-11-113-424-76	Sequence 76, App1	670	25	54.3	375	11	US-11-087-099-8969	Sequence 8969, App
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604	25	54.3	259	11	US-11-188-298-14615	Sequence 14615, A	677	25	54.3	384	11	US-11-087-099-327	Sequence 327, App
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716	25	54.3	492	11	US-11-264-096-1232	Sequence 1232, Ap	789	25	54.3	1003	11	US-11-204-755-11	Sequence 11, Appl			
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735	25	54.3	561	11	US-11-188-298-7184	Sequence 7184, Ap	808	25	54.3	1890	11	US-11-033-039-314	Sequence 314, App			
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857	24	51.3	24	11	US-11-004-399-1387	Sequence 1387, Ap	920	24	52.2	204	11	US-11-096-568A-20799	Sequence 20799, A
858	24	51.3	27	11	US-11-121-044A-90	Sequence 90, Appl	921	24	52.2	204	11	US-11-096-568A-28560	Sequence 28560, A
859	24	51.3	27	11	US-11-121-044A-124	Sequence 124, App	922	24	52.2	204	11	US-11-209-388-16	Sequence 16, Appl
860	24	51.3	31	11	US-11-004-399-167	Sequence 167, App	923	24	52.2	206	11	US-11-188-298-4493	Sequence 4493, Ap
861	24	51.3	42	9	US-10-957-887B-100	Sequence 100, App	924	24	52.2	208	9	US-10-467-657-4924	Sequence 4924, Ap
862	24	51.3	48	11	US-11-000-463-813	Sequence 341, App	925	24	52.2	212	11	US-11-123-441-108	Sequence 108, App
863	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	926	24	52.2	216	11	US-11-079-463-8530	Sequence 8530, Ap
864	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	927	24	52.2	216	11	US-11-079-463-8530	Sequence 10245, A
865	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	928	24	52.2	218	11	US-11-188-298-7023	Sequence 7023, Ap
866	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	929	24	52.2	218	11	US-11-045-004-390	Sequence 390, App
867	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	930	24	52.2	219	11	US-11-096-568A-6335	Sequence 6335, Ap
868	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	931	24	52.2	221	11	US-11-051-120-1776	Sequence 1776, Ap
869	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	932	24	52.2	221	11	US-11-043-806-574	Sequence 574, App
870	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	933	24	52.2	225	9	US-10-469-469-7	Sequence 7, Appl
871	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	934	24	52.2	225	11	US-11-156-516-38	Sequence 38, Appl
872	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	935	24	52.2	226	11	US-11-096-568A-29157	Sequence 29157, A
873	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	936	24	52.2	227	11	US-11-124-368A-231	Sequence 231, App
874	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	937	24	52.2	229	9	US-10-242-586-8	Sequence 8, Appl
875	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	938	24	52.2	229	9	US-10-242-902-8	Sequence 8, Appl
876	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	939	24	52.2	229	9	US-10-243-116-8	Sequence 8, Appl
877	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	940	24	52.2	229	9	US-10-243-116-8	Sequence 8, Appl
878	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	941	24	52.2	229	9	US-10-243-116-8	Sequence 8, Appl
879	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	942	24	52.2	229	9	US-10-243-218-8	Sequence 8, Appl
880	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	943	24	52.2	229	9	US-10-243-236-8	Sequence 8, Appl
881	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	944	24	52.2	229	9	US-10-243-236-8	Sequence 8, Appl
882	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	945	24	52.2	229	9	US-10-243-338-8	Sequence 8, Appl
883	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	946	24	52.2	229	9	US-10-243-348-8	Sequence 8, Appl
884	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	947	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
885	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	948	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
886	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	949	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
887	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	950	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
888	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	951	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
889	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	952	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
890	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	953	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
891	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	954	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
892	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	955	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
893	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	956	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
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895	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	958	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
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897	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	960	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
898	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	961	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
899	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	962	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
900	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	963	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
901	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	964	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
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903	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	966	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
904	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	967	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
905	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	968	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
906	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	969	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
907	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	970	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
908	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	971	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
909	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	972	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
910	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	973	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
911	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	974	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
912	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	975	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
913	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	976	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
914	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	977	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
915	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	978	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
916	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	979	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
917	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	980	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
918	24	51.3	48	11	US-11-000-463-813	Sequence 813, App	981	24	52.2	229	9	US-10-243-357-8	Sequence 8, Appl
919	24	51.3	48	11	US-11-000-4								

971 24 52.2 254 11 US-11-096-568A-28412 Sequence 28412, A
972 24 52.2 255 11 US-11-096-568A-29125 Sequence 29125, A
973 24 52.2 257 11 US-11-188-298-13446 Sequence 13446, A
974 24 52.2 258 11 US-11-188-298-11673 Sequence 11673, A
975 24 52.2 259 11 US-11-188-298-8953 Sequence 8953, A
976 24 52.2 261 9 US-10-528-031-6 Sequence 6, Appl
977 24 52.2 261 11 US-11-079-463-10157 Sequence 10157, A
978 24 52.2 262 9 US-10-793-626-2622 Sequence 2622, Ap
979 24 52.2 263 11 US-11-072-512-2517 Sequence 3517, Ap
980 24 52.2 267 11 US-11-096-568A-17567 Sequence 17567, A
981 24 52.2 269 11 US-11-188-298-16681 Sequence 16681, A
982 24 52.2 270 9 US-10-467-657-5806 Sequence 5806, A
983 24 52.2 273 11 US-11-188-298-9120 Sequence 9120, Ap
984 24 52.2 276 11 US-11-096-568A-28559 Sequence 28559, A
985 24 52.2 279 11 US-11-096-568A-33858 Sequence 33858, A
986 24 52.2 282 11 US-11-096-568A-15872 Sequence 15872, A
987 24 52.2 284 11 US-11-096-568A-29156 Sequence 29156, A
988 24 52.2 284 9 US-10-469-469-4 Sequence 4, Appl
989 24 52.2 286 11 US-11-188-298-17213 Sequence 17213, A
990 24 52.2 291 11 US-11-045-004-2015 Sequence 2015, Ap
991 24 52.2 297 11 US-11-188-298-13624 Sequence 13624, A
992 24 52.2 297 11 US-11-188-298-19279 Sequence 19279, A
993 24 52.2 299 9 US-10-467-657-2190 Sequence 2190, Ap
994 24 52.2 299 11 US-11-079-463-5793 Sequence 5793, Ap
995 24 52.2 301 9 US-10-877-346-73 Sequence 73, Appl
996 24 52.2 302 11 US-11-096-568A-10878 Sequence 10878, A
997 24 52.2 303 11 US-11-033-030-48 Sequence 48, Appl
998 24 52.2 304 11 US-11-096-568A-29277 Sequence 29277, A
999 24 52.2 304 11 US-11-033-030-47 Sequence 47, Appl
1000 24 52.2 307 9 US-10-469-469-21 Sequence 21, Appl

ALIGNMENTS

RESULT 1
US-10-530-253-13
Sequence 13, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casasetti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530, 253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415, 929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 13
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-13

Query Match 100.0%; Score 46; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
Db 15 LCTELQTTI 23

RESULT 2
US-11-206-138-3
Sequence 3, Application US/11206138
Publication No. US20060039919A1

GENERAL INFORMATION:
APPLICANT: Healthbanc Biotech CO, LTD.
TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
FILE REFERENCE: P7819/0613
CURRENT APPLICATION NUMBER: US/11/206,138
PRIOR FILING DATE: 2005-08-18
NUMBER OF SEQ ID NOS: 14
SOFTWARE: PatentIn version 3.3
SEQ ID NO 3
LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-11-206-138-3

Query Match 100.0%; Score 46; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.026;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
Db 22 LCTELQTTI 30

RESULT 3
US-10-530-253-1
Sequence 1, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casasetti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530, 253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415, 929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 1
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-1

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
| | | | |
Db 15 LCTELQTTI 23

RESULT 4
US-10-530-253-3
Sequence 3, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casasetti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530, 253
PRIOR FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415, 929

PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 3
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 5
US-10-530-253-5
Sequence 5, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 5
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 15 LCTELQTTI 23

RESULT 6
US-10-530-253-7
Sequence 7, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 7
LENGTH: 248

TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 112 LCTELQTTI 120

RESULT 7
US-10-530-253-9
Sequence 9, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 9
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LCTELQTTI 9
Db 112 LCTELQTTI 120

RESULT 8
US-10-530-253-11
Sequence 11, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 11
LENGTH: 248
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 46; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.04; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 LCTELOTTI 9
Db 112 LCTELOTTI 120

RESULT 9
US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; PRIOR FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 46; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.042; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 LCTELOTTI 9
Db 120 LCTELOTTI 128

RESULT 10
US-11-079-463-6325
; Sequence 6325, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTEROIDES FR
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; PRIOR FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6325
; LENGTH: 254
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-6325

Query Match 80.4%; Score 37; DB 11; Length 254;
Best Local Similarity 77.8%; Pred. No. 3; Indels 0; Gaps 0;
Matches 7; Conservative 1; Mismatches 1;

Qy 1 LCTELOTTI 9
Db 78 LCTELRVTI 86

RESULT 11
US-10-530-253-15

; Sequence 15, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 15
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 18
US-10-530-253-15

Query Match 76.1%; Score 35; DB 9; Length 158;
Best Local Similarity 66.7%; Pred. No. 4.8; Indels 0; Gaps 0;
Matches 6; Conservative 2; Mismatches 1;

Qy 1 LCTELOTTI 9
Db 17 LCTELNTSL 25

RESULT 12
US-10-530-253-20
; Sequence 20, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 45
US-10-530-253-20

Query Match 76.1%; Score 35; DB 9; Length 158;
Best Local Similarity 66.7%; Pred. No. 4.8; Indels 0; Gaps 0;
Matches 6; Conservative 2; Mismatches 1;

Qy 1 LCTELOTTI 9
Db 17 LCTELNTSL 25

RESULT 13
US-11-188-298-296
; Sequence 296, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT

FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 296
LENGTH: 587
TYPE: PRT
ORGANISM: Saccharomyces cerevisiae
US-11-188-298-296

Query Match 73.9%; Score 34; DB 11; Length 587;
Best Local Similarity 66.7%; Pred. No. 28;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 144 VCTILQTTV 152

RESULT 14
US-10-530-253-19
Sequence 19, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casareti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 19
LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 39
US-10-530-253-19

Query Match 71.7%; Score 33; DB 9; Length 158;
Best Local Similarity 66.7%; Pred. No. 12;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 17 LCTVLDTTL 25

RESULT 15
US-11-087-099-6055
Sequence 6055, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 6055
LENGTH: 420
TYPE: PRT
ORGANISM: Triticum aestivum
FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(420)

OTHER INFORMATION: unsure at all Xaa locations
US-11-087-099-6055

Query Match 71.7%; Score 33; DB 11; Length 420;
Best Local Similarity 65.7%; Pred. No. 33;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 CTELQTT 8
Db 308 CTERQTT 314

RESULT 16
US-11-096-568A-14735
Sequence 14735, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 14735
LENGTH: 492
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(492)
OTHER INFORMATION: Ceres Seq. ID no. 11050218
US-11-096-568A-14735

Query Match 71.7%; Score 33; DB 11; Length 492;
Best Local Similarity 77.8%; Pred. No. 38;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 43 LCTELATGI 51

RESULT 17
US-11-096-568A-14734
Sequence 14734, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 14734
LENGTH: 500
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(500)
OTHER INFORMATION: Ceres Seq. ID no. 11050217
US-11-096-568A-14734

Query Match 71.7%; Score 33; DB 11; Length 500;
Best Local Similarity 77.8%; Pred. No. 39;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 LCTELQTTI 9
Db 51 LCTELATGI 59

```
RESULT 18
US-11-096-568A-28552
; Sequence 28552, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 28552
; LENGTH: 1310
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(1310)
; OTHER INFORMATION: Ceres Seq. ID no. 2999574
US-11-096-568A-28552

Query Match          71.7%; Score 33; DB 11; Length 1310;
Best Local Similarity 62.5%; Pred. No. 1e+02; 0; Indels 0; Gaps 0;
Matches 5; Conservative 3; Mismatches 0;

QY      2 CTELOTTI 9
Db      953 CSELOTSV 960

RESULT 19
US-11-096-568A-28551
; Sequence 28551, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 28551
; LENGTH: 1368
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(1368)
; OTHER INFORMATION: Ceres Seq. ID no. 2999573
US-11-096-568A-28551

Query Match          71.7%; Score 33; DB 11; Length 1368;
Best Local Similarity 62.5%; Pred. No. 1e+02; 0; Indels 0; Gaps 0;
Matches 5; Conservative 3; Mismatches 0;

QY      2 CTELOTTI 9
Db      1011 CSELOTSV 1018

RESULT 20
US-11-096-568A-28550
; Sequence 28550, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
```

```
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 28550
; LENGTH: 1374
; TYPE: PRT
; ORGANISM: Arabidopsis thaliana
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(1374)
; OTHER INFORMATION: Ceres Seq. ID no. 2999572
US-11-096-568A-28550

Query Match          71.7%; Score 33; DB 11; Length 1374;
Best Local Similarity 62.5%; Pred. No. 1e+02; 0; Indels 0; Gaps 0;
Matches 5; Conservative 3; Mismatches 0;

QY      2 CTELOTTI 9
Db      1017 CSELOTSV 1024

RESULT 21
US-10-467-657-7248
; Sequence 7248, Application US/10467657
; Publication No. US20050260581A1
; GENERAL INFORMATION:
; APPLICANT: CHIRON SpA
; APPLICANT: FONTANA Maria Rita
; APPLICANT: PIZZA Mariagrazia
; APPLICANT: MIGINANI Vega
; APPLICANT: MONACI Elisabetta
; TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
; FILE REFERENCE:
; CURRENT APPLICATION NUMBER: US/10/467,657
; CURRENT FILING DATE: 2003-08-11
; PRIOR APPLICATION NUMBER: GB-0103424.8
; PRIOR FILING DATE: 2001-02-12
; NUMBER OF SEQ ID NOS: 9218
; SOFTWARE: SeqWIn99, version 1.04
; SEQ ID NO 7248
; LENGTH: 43
; TYPE: PRT
; ORGANISM: Neisseria gonorrhoeae
US-10-467-657-7248

Query Match          67.4%; Score 31; DB 9; Length 43;
Best Local Similarity 71.4%; Pred. No. 8.9;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      2 CTELOTT 8
Db      13 CTELEST 19

RESULT 22
US-10-519-390-15
; Sequence 15, Application US/10519390
; Publication No. US20060008872A1
; GENERAL INFORMATION:
; APPLICANT: MEDEXGEN Inc.
; APPLICANT: CHUNG, Yong-Hoon
; APPLICANT: LEE, Hak-sup
; APPLICANT: YI, Ki-Wan
; APPLICANT: KIM, Jae-Youn
; APPLICANT: HEO, Youn-Hwa
; TITLE OF INVENTION: A method of improving efficacy of biological response-modifying
; FILE REFERENCE: proteins and the example mutants
; CURRENT APPLICATION NUMBER: US/10/519,390
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: KR10-2003-0051846
; PRIOR FILING DATE: 2003-07-26
; NUMBER OF SEQ ID NOS: 65
```



```
; SOFTWARE: KopatentIn 1.71
; SEQ ID NO 15
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: IL-4: 33rd, 45th, 55th, 73rd, 82nd or 112nd Phe is replaced by
; OTHER INFORMATION: Val.
US-10-519-390-15
```

```
Query Match      67.4%; Score 31; DB 9; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 LCTELQTT 8
        |||||
Db      23 LCTELTWT 30
```

RESULT 23

```
US-11-176-830-207
; Sequence 207, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 207
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: Genbank AAA59149
; DATABASE ENTRY DATE: 1995-01-06
US-11-176-830-207
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 LCTELQTT 8
        |||||
Db      23 LCTELTWT 30
```

RESULT 24

```
US-11-176-830-546
; Sequence 546, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
```

```
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 546
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-546
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 LCTELQTT 8
        |||||
Db      23 LCTELTWT 30
```

RESULT 25

```
US-11-176-830-547
; Sequence 547, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 547
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-547
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 LCTELQTT 8
        |||||
Db      23 LCTELTWT 30
```

RESULT 26

```
US-11-176-830-548
; Sequence 548, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
```

```
;; CURRENT FILING DATE: 2005-07-06
;; PRIOR APPLICATION NUMBER: 10/658,834
;; PRIOR FILING DATE: 2003-09-08
;; PRIOR APPLICATION NUMBER: 60/457,135
;; PRIOR FILING DATE: 2003-03-21
;; PRIOR APPLICATION NUMBER: 60/409,898
;; PRIOR FILING DATE: 2002-09-09
;; NUMBER OF SEQ ID NOS: 1306
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO: 548
;; LENGTH: 129
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-11-176-830-548
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY      1 LCTELQTT 8
        |||||
Db      23 LCTELVT 30
```

```
RESULT 27
US-11-176-830-549
;; Sequence 549, Application US/11176830
;; Publication No. US20060020116A1
;; GENERAL INFORMATION:
;; APPLICANT: Gantier, Rene
;; APPLICANT: Guyon, Thierry
;; APPLICANT: Dittant, Lila
;; APPLICANT: Vega, Manuel
;; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
;; FILE REFERENCE: 17109-012002 (922B)
;; CURRENT APPLICATION NUMBER: US/11/176,830
;; CURRENT FILING DATE: 2005-07-06
;; PRIOR APPLICATION NUMBER: 10/658,834
;; PRIOR FILING DATE: 2003-09-08
;; PRIOR APPLICATION NUMBER: 60/457,135
;; PRIOR FILING DATE: 2003-03-21
;; PRIOR APPLICATION NUMBER: 60/409,898
;; PRIOR FILING DATE: 2002-09-09
;; NUMBER OF SEQ ID NOS: 1306
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO: 549
;; LENGTH: 129
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-11-176-830-549
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY      1 LCTELQTT 8
        |||||
Db      23 LCTELVT 30
```

```
RESULT 28
US-11-176-830-550
;; Sequence 550, Application US/11176830
;; Publication No. US20060020116A1
;; GENERAL INFORMATION:
;; APPLICANT: Gantier, Rene
;; APPLICANT: Guyon, Thierry
;; APPLICANT: Dittant, Lila
;; APPLICANT: Vega, Manuel
;; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
;; FILE REFERENCE: 17109-012002 (922B)
```

```
;; CURRENT APPLICATION NUMBER: US/11/176,830
;; CURRENT FILING DATE: 2005-07-06
;; PRIOR APPLICATION NUMBER: 10/658,834
;; PRIOR FILING DATE: 2003-09-08
;; PRIOR APPLICATION NUMBER: 60/457,135
;; PRIOR FILING DATE: 2003-03-21
;; PRIOR APPLICATION NUMBER: 60/409,898
;; PRIOR FILING DATE: 2002-09-09
;; NUMBER OF SEQ ID NOS: 1306
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO: 550
;; LENGTH: 129
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-11-176-830-550
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY      1 LCTELQTT 8
        |||||
Db      23 LCTELVT 30
```

```
RESULT 29
US-11-176-830-551
;; Sequence 551, Application US/11176830
;; Publication No. US20060020116A1
;; GENERAL INFORMATION:
;; APPLICANT: Gantier, Rene
;; APPLICANT: Guyon, Thierry
;; APPLICANT: Dittant, Lila
;; APPLICANT: Vega, Manuel
;; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
;; FILE REFERENCE: 17109-012002 (922B)
;; CURRENT APPLICATION NUMBER: US/11/176,830
;; CURRENT FILING DATE: 2005-07-06
;; PRIOR APPLICATION NUMBER: 10/658,834
;; PRIOR FILING DATE: 2003-09-08
;; PRIOR APPLICATION NUMBER: 60/457,135
;; PRIOR FILING DATE: 2003-03-21
;; PRIOR APPLICATION NUMBER: 60/409,898
;; PRIOR FILING DATE: 2002-09-09
;; NUMBER OF SEQ ID NOS: 1306
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO: 551
;; LENGTH: 129
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-11-176-830-551
```

```
Query Match      67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY      1 LCTELQTT 8
        |||||
Db      23 LCTELVT 30
```

```
RESULT 30
US-11-176-830-552
;; Sequence 552, Application US/11176830
;; Publication No. US20060020116A1
;; GENERAL INFORMATION:
;; APPLICANT: Gantier, Rene
;; APPLICANT: Guyon, Thierry
;; APPLICANT: Dittant, Lila
;; APPLICANT: Vega, Manuel
;; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
```

```
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 552
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-552
```

```
Query Match 67.4% Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
Db 23 LCTELTVT 30
```

RESULT 31

```
US-11-176-830-553
Sequence 553, Application US/11/176830
Publication No. US20060020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
APPLICANT: Vega, Manuel
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 553
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-553
```

```
Query Match 67.4% Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
Db 23 LCTELTVT 30
```

RESULT 32

```
US-11-176-830-554
Sequence 554, Application US/11/176830
Publication No. US20060020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
APPLICANT: Vega, Manuel
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
```

```
TITLE OF INVENTION: Acid Molecules and Related Applications
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 554
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-554
```

```
Query Match 67.4% Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
Db 23 LCTELTVT 30
```

RESULT 33

```
US-11-176-830-555
Sequence 555, Application US/11/176830
Publication No. US20060020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
APPLICANT: Vega, Manuel
TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
FILE REFERENCE: 17109-012002 (922B)
CURRENT APPLICATION NUMBER: US/11/176,830
CURRENT FILING DATE: 2005-07-06
PRIOR APPLICATION NUMBER: 10/658,834
PRIOR FILING DATE: 2003-09-08
PRIOR APPLICATION NUMBER: 60/457,135
PRIOR FILING DATE: 2003-03-21
PRIOR APPLICATION NUMBER: 60/409,898
PRIOR FILING DATE: 2002-09-09
NUMBER OF SEQ ID NOS: 1306
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 555
LENGTH: 129
TYPE: PRT
ORGANISM: Homo sapiens
US-11-176-830-555
```

```
Query Match 67.4% Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
Db 23 LCTELTVT 30
```

RESULT 34

```
US-11-176-830-556
Sequence 556, Application US/11/176830
Publication No. US20060020116A1
GENERAL INFORMATION:
APPLICANT: Gantier, Rene
APPLICANT: Guyon, Thierry
APPLICANT: Dittanti, Lila
APPLICANT: Vega, Manuel
```

```

; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 556
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-556
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELTYT 30
```

```

RESULT 35
US-11-176-830-557
; Sequence 557, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 557
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-557
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELTYT 30
```

```

RESULT 36
US-11-176-830-558
; Sequence 558, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlant, Lila
```

```

; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 558
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-558
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELTYT 30
```

```

RESULT 37
US-11-176-830-559
; Sequence 559, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirlant, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nuc
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 559
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-559
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELTYT 30
```

```

RESULT 38
US-11-176-830-560
; Sequence 560, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
```

```

; APPLICANT: Dirltanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; PRIOR FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 560
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-560
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```

Qy 1 LCTELQTT 8
    |||||
    23 LCTELVT 30
```

```

RESULT 39
US-11-176-830-561
; Sequence 561, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirltanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 561
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-561
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```

Qy 1 LCTELQTT 8
    |||||
    23 LCTELVT 30
```

```

RESULT 40
US-11-176-830-562
; Sequence 562, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
```

```

; APPLICANT: Guyon, Thierry
; APPLICANT: Dirltanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 562
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-562
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```

Qy 1 LCTELQTT 8
    |||||
    23 LCTELVT 30
```

```

RESULT 41
US-11-176-830-563
; Sequence 563, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dirltanti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cytokines for Higher Stability, Encoding Nu
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 563
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-563
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```

Qy 1 LCTELQTT 8
    |||||
    23 LCTELVT 30
```

```

RESULT 42
US-11-176-830-564
; Sequence 564, Application US/11176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
```

```

; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cyclokinines for Higher Stability, Encoding Nuc
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 564
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-564
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELVT 30
```

```

RESULT 43
US-11-176-830-565
; Sequence 565, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cyclokinines for Higher Stability, Encoding Nuc
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 565
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-565
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELVT 30
```

```

RESULT 44
US-11-176-830-566
; Sequence 566, Application US/11/176830
; Publication No. US20060020116A1
```

```

; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cyclokinines for Higher Stability, Encoding Nuc
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 566
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-566
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELVT 30
```

```

RESULT 45
US-11-176-830-567
; Sequence 567, Application US/11/176830
; Publication No. US20060020116A1
; GENERAL INFORMATION:
; APPLICANT: Gantier, Rene
; APPLICANT: Guyon, Thierry
; APPLICANT: Dittianti, Lila
; APPLICANT: Vega, Manuel
; TITLE OF INVENTION: Rational Evolution of Cyclokinines for Higher Stability, Encoding Nuc
; TITLE OF INVENTION: Acid Molecules and Related Applications
; FILE REFERENCE: 17109-012002 (922B)
; CURRENT APPLICATION NUMBER: US/11/176,830
; CURRENT FILING DATE: 2005-07-06
; PRIOR APPLICATION NUMBER: 10/658,834
; PRIOR FILING DATE: 2003-09-08
; PRIOR APPLICATION NUMBER: 60/457,135
; PRIOR FILING DATE: 2003-03-21
; PRIOR APPLICATION NUMBER: 60/409,898
; PRIOR FILING DATE: 2002-09-09
; NUMBER OF SEQ ID NOS: 1306
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 567
; LENGTH: 129
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-176-830-567
```

```

Query Match          67.4%; Score 31; DB 11; Length 129;
Best Local Similarity 75.0%; Pred. No. 26;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
    |||||
Db 23 LCTELVT 30
```

```

RESULT 46
US-10-530-253-17
; Sequence 17, Application US/10530253
```

```
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinny
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530.253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 17
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 33
US-10-530-253-17
```

```
Query Match          67.4%; Score 31; DB 9; Length 149;
Best Local Similarity 66.7%; Pred. No. 30;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTTI 9
    |||||
Db 15 LCGALETTI 23
```

```
RESULT 47
US-10-469-561-25
; Sequence 25, Application US/10469561
; Publication No. US20050260216A1
; GENERAL INFORMATION:
; APPLICANT: Claire Ashman
; APPLICANT: James Scott Crowe
; APPLICANT: Jonathan Henry Ellis
; APPLICANT: Alan Peter Lewis
; TITLE OF INVENTION: VACCINE
; FILE REFERENCE: PG4355USW
; CURRENT APPLICATION NUMBER: US/10/469,561
; CURRENT FILING DATE: 2003-08-29
; NUMBER OF SEQ ID NOS: 25
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 25
; LENGTH: 150
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric IL4 for human use
US-10-469-561-25
```

```
Query Match          67.4%; Score 31; DB 9; Length 150;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
    |||||
Db 48 LCTELTVT 55
```

```
RESULT 48
US-10-511-937-2476
; Sequence 2476, Application US/10511937
; Publication No. US20060088836A1
; GENERAL INFORMATION:
; APPLICANT: EXPRESSION DIAGNOSTICS, INC.
; APPLICANT: Wohlgenuth, Jay
; APPLICANT: Fry, Kirk
; APPLICANT: Woodward, Robert
; APPLICANT: Ly, Ngoc
```

```
; APPLICANT: Prentice, James
; APPLICANT: Morris, Macdonald
; APPLICANT: Rosenberg, Steven
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR DIAGNOSING
; TITLE OF INVENTION: AND MONITORING TRANSPLANT REJECTION
; FILE REFERENCE: 50661200104
; CURRENT APPLICATION NUMBER: US/10/511,937
; CURRENT FILING DATE: 2004-10-19
; PRIOR APPLICATION NUMBER: PCT/US2003/012946
; PRIOR FILING DATE: 2003-04-24
; PRIOR APPLICATION NUMBER: US 10/131,831
; PRIOR FILING DATE: 2002-04-24
; PRIOR APPLICATION NUMBER: US 10/325,899
; PRIOR FILING DATE: 2002-12-20
; NUMBER OF SEQ ID NOS: 3117
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2476
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-511-937-2476
```

```
Query Match          67.4%; Score 31; DB 8; Length 153;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
    |||||
Db 47 LCTELTVT 54
```

```
RESULT 49
US-11-174-398-8
; Sequence 8, Application US/11174398
; Publication No. US20050244930A1
; GENERAL INFORMATION:
; APPLICANT: Freeman, Scott R.
; APPLICANT: West, James W.
; APPLICANT: Novak, Julia E.
; TITLE OF INVENTION: ZALPHAL1 LIGAND ANTAGONISTS
; FILE REFERENCE: 01-37
; CURRENT APPLICATION NUMBER: US/11/174,398
; CURRENT FILING DATE: 2005-07-01
; PRIOR APPLICATION NUMBER: US/10/282,622
; PRIOR FILING DATE: 2002-10-28
; PRIOR APPLICATION NUMBER: 60/337,586
; PRIOR FILING DATE: 2001-11-05
; NUMBER OF SEQ ID NOS: 30
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Homo sapiens
US-11-174-398-8
```

```
Query Match          67.4%; Score 31; DB 11; Length 153;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
OY 1 LCTELQTT 8
    |||||
Db 47 LCTELTVT 54
```

```
RESULT 50
US-11-289-226-13
; Sequence 13, Application US/11289226
; Publication No. US20060084623A1
; GENERAL INFORMATION:
; APPLICANT: Steinman, Lawrence
; APPLICANT: Ruiz, Pedro
; APPLICANT: Garren, Hideki
; TITLE OF INVENTION: DNA Vaccination for Treatment of
```

```
; TITLE OF INVENTION: Autoimmune Disease
; FILE REFERENCE: STAN123CIP
; CURRENT APPLICATION NUMBER: US/11/289,226
; CURRENT FILING DATE: 2005-11-28
; PRIOR APPLICATION NUMBER: US/09/947,770
; PRIOR FILING DATE: 2001-09-05
; PRIOR APPLICATION NUMBER: PCT/US00/06233
; PRIOR FILING DATE: 2000-03-10
; PRIOR APPLICATION NUMBER: US 09/267,590
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: amino acid sequence of a Th2 cytokine
US-11-289-226-13
```

```
Query Match 67.4%; Score 31; DB 11; Length 153;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
Oy 1 LCTELQTT 8
Db 47 LCTELTVT 54
```

```
Search completed: May 5, 2006, 08:56:08
Job time : 10 secs
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GenCore version 5.1.7
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OM protein - protein search, using BW model

Run on: May 5, 2006, 02:25:57 ; Search time 20.7 Seconds
(without alignments)
35.946 Million cell updates/sec

Title: US-08-170-344-5

Perfect score: 46

Sequence: 1 ELQTTIHDI 9

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 1000 summaries

Database :

1: Issued Patents_Aa:*

2: /cgn2_6/ptodata/1/1aa/5-COMB.pep:*

3: /cgn2_6/ptodata/1/1aa/6-COMB.pep:*

4: /cgn2_6/ptodata/1/1aa/H-COMB.pep:*

5: /cgn2_6/ptodata/1/1aa/PTCNS-COMB.pep:*

6: /cgn2_6/ptodata/1/1aa/BACKL1est.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	46	100.0	14	1	US-07-909-122-3
2	46	100.0	30	1	US-08-363-586-4
3	46	100.0	30	2	US-09-980-523A-4
4	46	100.0	151	2	US-09-701-080C-18
5	46	100.0	158	2	US-09-980-523A-2
6	46	100.0	162	1	US-08-316-239B-3
7	46	100.0	162	1	US-08-316-239B-4
8	46	100.0	172	2	US-08-860-165-14
9	46	100.0	172	2	US-09-359-382-14
10	46	100.0	182	1	US-08-117-083-10
11	46	100.0	243	2	US-09-462-993-1
12	46	100.0	266	2	US-08-860-165-10
13	46	100.0	266	2	US-09-359-382-10
14	46	100.0	266	2	US-09-359-382-10
15	46	100.0	273	2	US-09-485-885-4
16	46	100.0	292	2	US-09-485-885-10
17	46	100.0	371	2	US-09-485-885-10
18	46	100.0	390	2	US-09-485-885-14
19	41	89.1	9	1	US-08-787-547-102
20	41	89.1	20	1	US-08-159-339A-248
21	41	89.1	9	1	US-08-934-915-159
22	36	76.3	317	2	US-08-913-159-14
23	35	76.1	248	2	US-09-600-932-28
24	35	76.1	248	2	US-09-949-016-6612
25	35	76.1	251	2	US-09-949-016-8481
26	35	76.1	251	2	US-09-949-016-9058
27	35	76.1	259	2	US-09-949-016-9059

28	34	73.9	56	2	US-08-936-165A-297	Sequence 297, App
29	34	73.9	383	2	US-09-603-208A-62	Sequence 62, Appl
30	34	73.9	680	2	US-09-949-016-7565	Sequence 7565, Ap
31	34	73.9	695	2	US-09-248-796A-20895	Sequence 20895, A
32	34	73.9	811	2	US-09-252-991A-22216	Sequence 22216, A
33	34	73.9	1896	2	US-09-964-956-13	Sequence 13, Appl
34	34	73.9	1905	2	US-09-964-956-44	Sequence 44, Appl
35	33	71.7	59	1	US-08-118-270-256	Sequence 256, App
36	33	71.7	59	4	PCT-US93-08528-256	Sequence 256, App
37	33	71.7	166	2	US-09-270-767-33046	Sequence 33046, A
38	33	71.7	166	2	US-09-270-767-48263	Sequence 48263, A
39	33	71.7	198	2	US-09-134-001C-4050	Sequence 4050, Ap
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41	33	71.7	500	2	US-10-214-269-19	Sequence 19, Appl
42	33	71.7	503	2	US-09-990-337-3	Sequence 3, Appl
43	33	71.7	738	2	US-09-543-681A-7528	Sequence 7528, Ap
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45	33	71.7	1128	2	US-09-252-991A-31032	Sequence 31032, A
46	33	71.7	1769	2	US-09-949-016-8280	Sequence 8280, Ap
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56	32	69.6	513	2	US-09-769-787-193	Sequence 37, Appl
57	32	69.6	513	2	US-09-769-787-37	Sequence 37, Appl
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59	32	69.6	572	2	US-09-545-916-7	Sequence 6825, Ap
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61	31	67.4	336	2	US-09-252-991A-27158	Sequence 27158, A
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66	31	67.4	846	2	US-08-938-085A-33	Sequence 33, Appl
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69	31	67.4	846	2	US-10-072-841A-33	Sequence 33, Appl
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71	31	67.4	861	2	US-08-538-092-809	Sequence 809, App
72	31	67.4	1016	2	US-09-949-016-11304	Sequence 11304, A
73	31	67.4	1193	2	US-09-227-725A-4	Sequence 4, Appl
74	31	67.4	1193	2	US-10-071-900-4	Sequence 4, Appl
75	31	67.4	1211	2	US-09-167-206-14	Sequence 14, Appl
76	30	65.2	71	2	US-09-513-999C-5529	Sequence 5529, Ap
77	30	65.2	112	2	US-09-893-737-322	Sequence 322, App
78	30	65.2	136	2	US-09-303-518D-74	Sequence 74, Appl
79	30	65.2	183	2	US-09-252-991A-26451	Sequence 26451, A
80	30	65.2	198	2	US-09-540-226-3042	Sequence 3042, Ap
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82	30	65.2	229	2	US-09-303-518D-76	Sequence 76, Appl
83	30	65.2	230	2	US-09-252-991A-22090	Sequence 22090, A
84	30	65.2	230	2	US-09-270-767-43528	Sequence 43529, A
85	30	65.2	250	2	US-09-252-991A-21053	Sequence 21053, A
86	30	65.2	334	2	US-09-543-661A-6758	Sequence 6758, Ap
87	30	65.2	362	2	US-09-252-991A-31349	Sequence 31349, A
88	30	65.2	408	2	US-09-540-226-3813	Sequence 3813, Ap
89	30	65.2	408	2	US-09-489-029A-13552	Sequence 13552, A
90	30	65.2	414	2	US-09-605-703B-908	Sequence 908, App
91	30	65.2	419	1	US-08-686-417-3	Sequence 3, Appl
92	30	65.2	455	1	US-09-328-332-6800	Sequence 6800, Ap
93	30	65.2	455	1	US-08-819-013-1	Sequence 1, Appl
94	30	65.2	456	2	US-09-355-214-1	Sequence 5, Appl
95	30	65.2	456	2	US-09-355-214-5	Sequence 3013, Ap
96	30	65.2	469	2	US-09-107-433-3013	Sequence 7976, A
97	30	65.2	470	2	US-09-949-016-7976	Sequence 18644, A
98	30	65.2	488	2	US-09-248-796A-18644	Sequence 8155, Ap
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101	30	65.2	567	2	US-09-198-452A-994	Sequence 994, App	174	28	60.9	74	2	US-09-134-000C-6520	Sequence 6520, App
102	30	65.2	664	2	US-09-605-703B-906	Sequence 906, App	175	28	60.9	102	1	US-07-901-703-6	Sequence 6, App1
103	30	65.2	760	2	US-09-107-532A-5094	Sequence 5094, App	176	28	60.9	102	1	US-08-278-928A-12	Sequence 12, App1
104	30	65.2	745	2	US-09-177-165A-25	Sequence 25, App1	177	28	60.9	102	1	US-08-155-343A-12	Sequence 12, App1
105	30	65.2	909	2	US-09-902-540-12961	Sequence 12961, A	178	28	60.9	102	1	US-08-406-677-12	Sequence 12, App1
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108	30	65.2	1281	1	US-09-636-728-5	Sequence 5, App1	181	28	60.9	102	1	US-08-643-763A-12	Sequence 12, App1
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110	30	65.2	1298	1	US-08-843-530B-4	Sequence 4, App1	183	28	60.9	102	1	US-08-451-953A-12	Sequence 12, App1
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112	30	65.2	1298	1	US-09-636-728-2	Sequence 2, App1	185	28	60.9	102	1	US-08-461-397A-12	Sequence 12, App1
113	30	65.2	1881	2	US-09-233-086-3	Sequence 3, App1	186	28	60.9	102	1	US-08-912-088-12	Sequence 12, App1
114	30	65.2	3696	2	US-09-134-001C-5080	Sequence 5080, App	187	28	60.9	102	2	US-08-478-730A-12	Sequence 12, App1
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116	29	63.0	91	2	US-09-248-796A-25740	Sequence 25740, A	189	28	60.9	102	2	US-08-445-467-12	Sequence 12, App1
117	29	63.0	101	2	US-09-252-991A-30506	Sequence 30506, A	190	28	60.9	102	2	US-08-445-467-12	Sequence 12, App1
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119	29	63.0	140	2	US-09-640-111A-692	Sequence 692, App	192	28	60.9	102	2	US-08-414-033A-12	Sequence 12, App1
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121	29	63.0	168	2	US-09-710-379-2528	Sequence 2528, App	194	28	60.9	102	2	US-08-931-858E-165	Sequence 165, App
122	29	63.0	177	2	US-09-732-210-1075	Sequence 1075, App	195	28	60.9	102	2	US-08-981-739-165	Sequence 165, App
123	29	63.0	222	2	US-09-252-991A-20591	Sequence 20591, A	196	28	60.9	102	2	US-08-440-894A-12	Sequence 12, App1
124	29	63.0	237	2	US-09-134-001C-4185	Sequence 4185, App	197	28	60.9	102	2	US-09-170-936-12	Sequence 12, App1
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126	29	63.0	289	2	US-09-068-655-13	Sequence 13, App1	199	28	60.9	102	2	US-09-128-026-165	Sequence 165, App
127	29	63.0	290	2	US-09-068-655-9	Sequence 9, App1	200	28	60.9	102	2	US-09-496-398-7	Sequence 7, App1
128	29	63.0	291	2	US-09-107-532A-5063	Sequence 5063, App	201	28	60.9	102	2	US-09-496-398-8	Sequence 8, App1
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130	29	63.0	297	2	US-09-097-231-8	Sequence 8, App1	203	28	60.9	102	2	US-08-643-321-11	Sequence 11, App1
131	29	63.0	297	2	US-09-353-099-8	Sequence 8, App1	204	28	60.9	102	2	US-09-464-206-12	Sequence 12, App1
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134	29	63.0	334	2	US-09-674-826B-6	Sequence 6, App1	207	28	60.9	102	2	US-09-374-958C-46	Sequence 46, App1
135	29	63.0	335	1	US-08-903-800A-6	Sequence 6, App1	208	28	60.9	102	2	US-09-320-527-165	Sequence 165, App
136	29	63.0	337	2	US-09-328-352-5664	Sequence 5664, App	209	28	60.9	102	2	US-09-820-407-165	Sequence 165, App
137	29	63.0	345	2	US-10-200-012-20	Sequence 20, App1	210	28	60.9	102	2	US-09-513-999C-8150	Sequence 8150, App
138	29	63.0	402	2	US-09-540-236-2685	Sequence 2685, App	211	28	60.9	102	2	US-08-260-675-12	Sequence 12, App1
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140	29	63.0	437	1	US-08-989-925-3	Sequence 3, App1	213	28	60.9	102	2	US-08-292-782-12	Sequence 12, App1
141	29	63.0	437	1	US-09-949-016-9609	Sequence 9609, App	214	28	60.9	102	4	PCT-US92-01968-12	Sequence 12, App1
142	29	63.0	485	2	US-09-489-039A-10167	Sequence 10167, A	215	28	60.9	102	4	PCT-US93-05446-6	Sequence 6, App1
143	29	63.0	485	2	US-10-130-419-1	Sequence 1, App1	216	28	60.9	102	4	PCT-US93-07110-12	Sequence 12, App1
144	29	63.0	532	2	US-09-252-991A-30492	Sequence 30492, A	217	28	60.9	102	4	PCT-US93-07231-12	Sequence 12, App1
145	29	63.0	532	2	US-09-252-991A-30492	Sequence 30492, A	218	28	60.9	102	4	PCT-US93-08742-12	Sequence 12, App1
146	29	63.0	538	2	US-09-487-558B-200	Sequence 200, App	219	28	60.9	102	4	PCT-US93-08808-12	Sequence 12, App1
147	29	63.0	626	2	US-10-104-047-2449	Sequence 2449, App	220	28	60.9	102	4	PCT-US93-08808-12	Sequence 12, App1
148	29	63.0	672	2	US-09-487-558B-200	Sequence 200, App	221	28	60.9	110	2	US-09-902-540-13445	Sequence 13445, A
149	29	63.0	821	1	US-07-928-464-2	Sequence 2, App1	222	28	60.9	118	1	US-08-481-377-10	Sequence 10, App1
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152	29	63.0	850	4	PCT-US93-07347-2	Sequence 2, App1	225	28	60.9	118	2	US-08-946-092A-8	Sequence 8, App1
153	29	63.0	937	2	US-09-005-180A-4	Sequence 4, App1	226	28	60.9	118	2	US-09-172-062-8	Sequence 8, App1
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155	29	63.0	944	2	US-09-538-092-452	Sequence 452, App	228	28	60.9	118	2	US-09-389-705-10	Sequence 10, App1
156	29	63.0	988	2	US-09-540-236-3801	Sequence 3801, App	229	28	60.9	118	4	PCT-US94-00666-10	Sequence 10, App1
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161	29	63.0	3092	4	US-09-487-558B-172	Sequence 172, App	234	28	60.9	128	1	US-08-455-550-16	Sequence 16, App1
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165	28	60.9	33	1	US-09-557-034-75	Sequence 75, App1	238	28	60.9	139	2	US-09-270-767-43863	Sequence 43863, App
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169	28	60.9	60	2	US-08-744-231-27	Sequence 27, App1	242	28	60.9	187	2	US-09-270-767-37546	Sequence 37546, App
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172	28	60.9	72	2	US-09-270-767-52246	Sequence 52246, App	245	28	60.9	212	2	US-09-902-540-15176	Sequence 15176, App
173	28	60.9	73	2	US-09-107-532A-4579	Sequence 4579, App	246	28	60.9	224	1	US-08-173-510B-87	Sequence 87, App1

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250	28	60.9	224	2	US-08-060-433C-37	Sequence 37, Appl	323	28	60.9	466	2	US-09-635-504-35	Sequence 35, Appl
251	28	60.9	224	2	US-08-450-482B-87	Sequence 87, Appl	324	28	60.9	467	2	US-09-543-681A-7427	Sequence 7427, Ap
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253	28	60.9	224	2	US-09-248-796A-17942	Sequence 17942, A	326	28	60.9	482	2	US-09-107-433-4705	Sequence 4705, Ap
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257	28	60.9	287	2	US-09-489-039A-8878	Sequence 8878, Ap	330	28	60.9	520	2	US-09-328-352-7451	Sequence 8381, Ap
258	28	60.9	290	2	US-09-134-000C-5874	Sequence 5874, Ap	331	28	60.9	550	2	US-10-104-047-3416	Sequence 3416, Ap
259	28	60.9	302	1	US-09-031-485-38	Sequence 38, Appl	332	28	60.9	550	2	US-09-489-039A-8381	Sequence 19777, A
260	28	60.9	302	1	US-08-847-429A-38	Sequence 38, Appl	333	28	60.9	558	2	US-09-248-796A-19777	Sequence 14260, A
261	28	60.9	302	2	US-09-065-474-38	Sequence 38, Appl	334	28	60.9	547	2	US-09-134-000C-1260	Sequence 5126, Ap
262	28	60.9	302	2	US-09-457-046B-18	Sequence 18, Appl	335	28	60.9	617	2	US-09-344-000C-5126	Sequence 19796, A
263	28	60.9	302	2	US-09-557-034-38	Sequence 18, Appl	336	28	60.9	630	2	US-09-552-991A-19796	Sequence 5104, Ap
264	28	60.9	302	2	US-09-866-570B-18	Sequence 18, Appl	337	28	60.9	630	2	US-09-543-681A-5104	Sequence 4829, Ap
265	28	60.9	303	1	US-09-031-485-23	Sequence 23, Appl	338	28	60.9	650	2	US-08-674-351-2	Sequence 2, Appl
266	28	60.9	303	1	US-08-847-429A-23	Sequence 23, Appl	339	28	60.9	660	2	US-09-107-532A-6403	Sequence 6403, Ap
267	28	60.9	303	2	US-09-065-474-23	Sequence 23, Appl	340	28	60.9	688	2	US-09-489-039A-7813	Sequence 9613, Ap
268	28	60.9	303	2	US-09-557-034-23	Sequence 23, Appl	341	28	60.9	728	2	US-09-248-796A-16666	Sequence 16666, A
269	28	60.9	312	2	US-09-489-039A-10591	Sequence 10591, A	342	28	60.9	745	1	US-08-887-518-3	Sequence 3, Appl
270	28	60.9	319	2	US-10-028-051A-8	Sequence 8, Appl	343	28	60.9	745	1	US-09-023-321-3	Sequence 3, Appl
271	28	60.9	325	2	US-08-878-474-9	Sequence 9, Appl	344	28	60.9	745	1	US-08-890-853-4	Sequence 4, Appl
272	28	60.9	325	2	US-09-976-594-479	Sequence 479, App	345	28	60.9	745	1	US-09-032-475-3	Sequence 3, Appl
273	28	60.9	325	2	US-10-014-055-2	Sequence 2, Appl	346	28	60.9	745	1	US-09-099-125A-4	Sequence 4, Appl
274	28	60.9	325	2	US-10-014-055-4	Sequence 4, Appl	347	28	60.9	745	1	US-09-099-124A-4	Sequence 4, Appl
275	28	60.9	325	2	US-10-028-051A-2	Sequence 2, Appl	348	28	60.9	745	1	US-09-032-476-4	Sequence 4, Appl
276	28	60.9	325	2	US-10-028-051A-4	Sequence 4, Appl	349	28	60.9	745	2	US-08-890-854-4	Sequence 4, Appl
277	28	60.9	335	2	US-09-602-787A-230	Sequence 230, App	350	28	60.9	745	2	US-09-023-324-4	Sequence 4, Appl
278	28	60.9	343	2	US-09-270-767-39334	Sequence 39334, A	351	28	60.9	745	2	US-09-168-829-2	Sequence 2, Appl
279	28	60.9	343	2	US-09-270-767-54551	Sequence 54551, A	352	28	60.9	745	2	US-08-810-820-10	Sequence 10, Appl
280	28	60.9	351	1	US-08-929-417-2	Sequence 2, Appl	353	28	60.9	745	2	US-09-109-986-4	Sequence 2, Appl
281	28	60.9	352	2	US-09-065-474-139	Sequence 139, App	354	28	60.9	745	2	US-09-844-908-10	Sequence 3, Appl
282	28	60.9	352	2	US-09-557-034-139	Sequence 139, App	355	28	60.9	745	2	US-09-796-872-2	Sequence 2, Appl
283	28	60.9	352	2	US-09-198-452A-582	Sequence 382, App	356	28	60.9	745	2	US-09-248-796A-19137	Sequence 19137, A
284	28	60.9	353	2	US-09-618-912-204	Sequence 204, App	357	28	60.9	745	2	US-09-543-681A-4312	Sequence 4312, Ap
285	28	60.9	354	2	US-09-538-092-1227	Sequence 1227, Ap	358	28	60.9	745	2	US-09-252-991A-32826	Sequence 32826, A
286	28	60.9	354	2	US-09-248-796A-24790	Sequence 24790, A	359	28	60.9	745	2	US-09-328-352-5208	Sequence 5208, A
287	28	60.9	358	1	US-08-700-186-2	Sequence 2, Appl	360	28	60.9	745	2	US-09-417-197-123	Sequence 123, App
288	28	60.9	358	1	US-08-914-981-2	Sequence 2, Appl	361	28	60.9	745	2	US-09-417-197-121	Sequence 121, App
289	28	60.9	358	2	US-09-116-115-2	Sequence 2, Appl	362	28	60.9	745	2	US-09-139-802-201	Sequence 201, App
290	28	60.9	358	2	US-09-541-762-2	Sequence 546, App	363	28	60.9	745	2	US-09-659-786-201	Sequence 8, Appl
291	28	60.9	358	2	US-09-438-185A-546	Sequence 7301, Ap	364	28	60.9	745	2	US-09-646-533D-8	Sequence 123, App
292	28	60.9	359	2	US-09-949-016-7301	Sequence 13, Appl	365	28	60.9	745	2	US-09-417-197-123	Sequence 121, App
293	28	60.9	360	1	US-08-459-346-13	Sequence 4, Appl	366	28	60.9	745	2	US-09-417-197-121	Sequence 121, App
294	28	60.9	360	1	US-08-411-607A-4	Sequence 13, Appl	367	28	60.9	745	2	US-09-489-039A-14025	Sequence 31, Appl
295	28	60.9	360	2	US-08-889-419-13	Sequence 13, Appl	368	28	60.9	745	2	US-10-144-198-31	Sequence 31, Appl
296	28	60.9	360	2	US-08-402-542-13	Sequence 13, Appl	369	28	60.9	745	2	US-09-031-485-33	Sequence 33, Appl
297	28	60.9	360	2	US-09-361-741-4	Sequence 4, Appl	370	28	60.9	745	2	US-08-847-429A-33	Sequence 33, Appl
298	28	60.9	360	2	US-09-461-418-4	Sequence 13, Appl	371	28	60.9	745	2	US-09-065-474-33	Sequence 33, Appl
299	28	60.9	360	4	PCT-US93-07189-13	Sequence 7544, Ap	372	28	60.9	745	2	US-09-557-034-33	Sequence 6, Appl
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302	28	60.9	386	2	US-09-190-911-1	Sequence 11, Appl	375	28	60.9	745	2	US-09-045-201A-2	Sequence 2, Appl
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304	28	60.9	386	2	US-10-007-262-1	Sequence 2, Appl	377	28	60.9	745	2	US-09-137-484-72	Sequence 72, Appl
305	28	60.9	386	2	US-09-645-078-2	Sequence 4367, Ap	378	28	60.9	745	2	US-09-601-729-77	Sequence 72, Appl
306	28	60.9	388	2	US-09-543-681A-4387	Sequence 17335, A	379	28	60.9	745	2	FCT-US95-02121-72	Sequence 72, Appl
307	28	60.9	389	2	US-09-248-796A-17335	Sequence 15573, A	380	28	60.9	745	2	US-09-053-485-15	Sequence 35, Appl
308	28	60.9	414	2	US-09-248-796A-15573	Sequence 6642, Ap	381	28	60.9	745	2	US-08-466-285-2	Sequence 2, Appl
309	28	60.9	416	2	US-09-107-532A-6642	Sequence 7435, Ap	382	27	58.7	745	2	US-09-164-768-2	Sequence 2, Appl
310	28	60.9	418	2	US-09-543-681A-7435	Sequence 2, Appl	383	27	58.7	745	2	US-09-350-027-6	Sequence 6, Appl
311	28	60.9	438	2	US-09-044-718-2	Sequence 2, Appl	384	27	58.7	745	2	US-09-621-976-7086	Sequence 7086, Ap
312	28	60.9	438	2	US-10-062-848-2	Sequence 102, App	385	27	58.7	745	2	US-09-270-767-3481	Sequence 43481, A
313	28	60.9	440	2	US-09-684-855-102	Sequence 125, App	386	27	58.7	745	2	US-09-270-767-58842	Sequence 58842, A
314	28	60.9	440	2	US-09-684-855-125	Sequence 147, App	387	27	58.7	745	2	US-09-107-532A-6403	Sequence 6074, Ap
315	28	60.9	440	2	US-09-684-855-147	Sequence 2, Appl	388	27	58.7	745	2		
316	28	60.9	440	2	US-09-488-265B-2	Sequence 15, Appl	389	27	58.7	745	2		
317	28	60.9	450	2	US-09-044-718-15	Sequence 4333, Ap	390	27	58.7	745	2		
318	28	60.9	450	2	US-09-583-110-4333	Sequence 15, Appl	391	27	58.7	745	2		
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395	27	58.7	135	2	US-09-270-767-54860	Sequence 54860, A	468	27	58.7	395	2	US-09-949-016-11110	Sequence 11110, A
396	27	58.7	127	2	US-09-248-796A-18260	Sequence 18260, A	469	27	58.7	401	2	US-09-252-991A-27198	Sequence 27198, A
397	27	58.7	139	2	US-09-513-999C-4772	Sequence 4772, Ap	470	27	58.7	401	2	US-09-248-796A-17808	Sequence 17808, A
398	27	58.7	133	2	US-09-605-703B-790	Sequence 790, App	471	27	58.7	414	2	US-09-540-236-2868	Sequence 2868, Ap
399	27	58.7	147	2	US-09-605-703B-788	Sequence 788, App	472	27	58.7	415	2	US-09-949-016-11269	Sequence 11269, A
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401	27	58.7	154	2	US-09-345-336B-144	Sequence 144, App	474	27	58.7	426	2	US-08-961-084-48	Sequence 48, App1
402	27	58.7	158	1	US-08-247-504B-10	Sequence 10, App1	475	27	58.7	426	2	US-09-536-784-48	Sequence 48, App1
403	27	58.7	158	1	US-08-767-942A-19	Sequence 19, App1	476	27	58.7	426	2	US-09-765-271-48	Sequence 48, App1
404	27	58.7	163	2	US-09-053-197A-22	Sequence 22, App1	477	27	58.7	426	2	US-09-765-272A-48	Sequence 48, App1
405	27	58.7	153	2	US-09-085-761A-22	Sequence 22, App1	478	27	58.7	427	2	US-09-196-857-2	Sequence 2, App1
406	27	58.7	176	2	US-09-134-000C-4892	Sequence 4892, Ap	479	27	58.7	427	2	US-09-583-110-3940	Sequence 3940, Ap
407	27	58.7	181	2	US-09-248-796A-20509	Sequence 20509, A	480	27	58.7	435	2	US-09-107-433-4256	Sequence 4256, Ap
408	27	58.7	185	2	US-09-540-236-2830	Sequence 2830, Ap	481	27	58.7	442	2	US-09-248-796A-20425	Sequence 20425, A
409	27	58.7	187	2	US-09-134-001C-3251	Sequence 3251, Ap	482	27	58.7	449	1	US-08-924-663A-2	Sequence 2, App1
410	27	58.7	196	2	US-10-241-602B-4	Sequence 4, App1	483	27	58.7	449	1	US-08-974-565C-1	Sequence 1, App1
411	27	58.7	200	2	US-09-270-767-42004	Sequence 42004, A	484	27	58.7	449	2	US-09-255-748-1	Sequence 1, App1
412	27	58.7	214	2	US-09-238-986-129	Sequence 129, App	485	27	58.7	460	2	US-09-013-118-3	Sequence 3, App1
413	27	58.7	214	2	US-10-101-464A-129	Sequence 129, App	486	27	58.7	464	2	US-09-370-767-44193	Sequence 44193, A
414	27	58.7	216	2	US-09-345-336B-115	Sequence 115, App	487	27	58.7	466	2	US-09-489-039A-9530	Sequence 9530, Ap
415	27	58.7	216	2	US-09-489-039A-10545	Sequence 10545, A	488	27	58.7	476	2	US-09-949-016-10314	Sequence 10314, A
416	27	58.7	238	2	US-09-248-796A-14524	Sequence 14524, A	489	27	58.7	478	2	US-08-740-223A-7	Sequence 7, App1
417	27	58.7	249	2	US-09-248-796A-22159	Sequence 22159, A	490	27	58.7	478	2	US-09-709-188-7	Sequence 7, App1
418	27	58.7	254	2	US-09-248-796A-22744	Sequence 22744, A	491	27	58.7	478	2	US-10-225-060-7	Sequence 7, App1
419	27	58.7	258	2	US-10-101-464A-723	Sequence 723, App	492	27	58.7	479	2	US-08-985-343-4	Sequence 4, App1
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422	27	58.7	270	2	US-09-248-796A-14842	Sequence 14842, A	495	27	58.7	490	2	US-09-709-188-12	Sequence 12, App1
423	27	58.7	271	1	US-08-117-083-14	Sequence 14, App1	496	27	58.7	492	2	US-10-225-060-12	Sequence 12, App1
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425	27	58.7	278	2	US-09-485-885-21	Sequence 21, App1	498	27	58.7	495	2	US-08-740-223A-26	Sequence 26, App1
426	27	58.7	282	2	US-09-605-703B-1024	Sequence 1024, Ap	499	27	58.7	495	2	US-09-351-457-5	Sequence 5, App1
427	27	58.7	282	2	US-09-605-703B-1026	Sequence 1026, Ap	500	27	58.7	495	2	US-09-561-500-5	Sequence 5, App1
428	27	58.7	283	2	US-09-328-352-7680	Sequence 7680, Ap	501	27	58.7	495	2	US-09-561-108-5	Sequence 5, App1
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432	27	58.7	305	2	US-09-339-159B-18	Sequence 18, App1	505	27	58.7	495	2	US-09-561-499-5	Sequence 5, App1
433	27	58.7	305	2	US-09-107-532A-6585	Sequence 6585, Ap	506	27	58.7	495	2	US-09-998-803-5	Sequence 5, App1
434	27	58.7	308	2	US-09-252-991A-20337	Sequence 20337, A	507	27	58.7	495	2	US-09-561-001-5	Sequence 5, App1
435	27	58.7	311	2	US-09-878-781-18	Sequence 18, App1	508	27	58.7	495	2	US-09-819-386-5	Sequence 5, App1
436	27	58.7	313	2	US-09-876-216-2	Sequence 2, App1	509	27	58.7	495	2	US-09-351-598-5	Sequence 5, App1
437	27	58.7	313	2	US-10-359-076-2	Sequence 2, App1	510	27	58.7	495	2	US-10-225-060-26	Sequence 26, App1
438	27	58.7	318	2	US-09-107-532A-5824	Sequence 5824, Ap	511	27	58.7	495	2	US-09-562-245-5	Sequence 5, App1
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441	27	58.7	328	2	US-09-636-060C-8	Sequence 8, App1	514	27	58.7	497	1	US-08-665-926-4	Sequence 4, App1
442	27	58.7	328	2	US-09-986-552-8	Sequence 8, App1	515	27	58.7	497	1	US-08-348-492-4	Sequence 4, App1
443	27	58.7	328	2	US-09-636-596C-8	Sequence 8, App1	516	27	58.7	497	2	US-09-162-437-4	Sequence 4, App1
444	27	58.7	328	2	US-10-023-894-10	Sequence 10, App1	517	27	58.7	497	2	US-08-740-223A-4	Sequence 4, App1
445	27	58.7	328	2	US-10-306-686-8	Sequence 8, App1	518	27	58.7	497	2	US-08-740-223A-14	Sequence 14, App1
446	27	58.7	328	2	US-09-895-072-8	Sequence 8, App1	519	27	58.7	497	2	US-08-817-318-4	Sequence 4, App1
447	27	58.7	328	2	US-10-023-888-10	Sequence 10, App1	520	27	58.7	497	2	US-09-709-188-4	Sequence 4, App1
448	27	58.7	333	2	US-09-538-092-778	Sequence 778, App	521	27	58.7	497	2	US-09-709-188-14	Sequence 14, App1
449	27	58.7	335	2	US-09-949-016-6163	Sequence 6163, Ap	522	27	58.7	497	2	US-09-442-717-4	Sequence 4, App1
450	27	58.7	339	2	US-09-543-681A-4731	Sequence 4731, Ap	523	27	58.7	497	2	US-09-689-020-4	Sequence 4, App1
451	27	58.7	341	2	US-09-543-681A-8334	Sequence 8334, Ap	524	27	58.7	497	2	US-10-225-060-4	Sequence 4, App1
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454	27	58.7	342	2	US-09-252-991A-32983	Sequence 32983, A	527	27	58.7	498	1	US-08-418-595-2	Sequence 2, App1
455	27	58.7	350	2	US-09-543-681A-4618	Sequence 4618, Ap	528	27	58.7	498	1	US-08-665-926-2	Sequence 2, App1
456	27	58.7	360	2	US-09-949-016-8354	Sequence 8354, Ap	529	27	58.7	498	1	US-08-348-492-2	Sequence 2, App1
457	27	58.7	365	1	US-09-068-109-2	Sequence 2, App1	530	27	58.7	498	2	US-09-162-437-2	Sequence 2, App1
458	27	58.7	365	2	US-09-489-847-322	Sequence 322, App	531	27	58.7	498	2	US-08-740-223A-2	Sequence 2, App1
459	27	58.7	369	2	US-09-248-796A-18353	Sequence 18353, A	532	27	58.7	498	2	US-08-740-223A-20	Sequence 20, App1
460	27	58.7	369	2	US-09-543-681A-7173	Sequence 7173, Ap	533	27	58.7	498	2	US-09-351-457-2	Sequence 2, App1
461	27	58.7	376	2	US-09-198-452A-1112	Sequence 1112, Ap	534	27	58.7	498	2	US-09-561-108-2	Sequence 2, App1
462	27	58.7	383	2	US-09-485-885-23	Sequence 23, App1	535	27	58.7	498	2	US-09-561-543-2	Sequence 2, App1
463	27	58.7	388	2	US-09-438-185A-1038	Sequence 1038, Ap	536	27	58.7	498	2	US-09-561-526-2	Sequence 2, App1
464	27	58.7	391	2	US-09-270-767-43901	Sequence 43901, A	537	27	58.7	498	2	US-09-561-526-2	Sequence 2, App1
465	27	58.7	391	2	US-09-987-107-38	Sequence 38, App1	538	27	58.7	498	2	US-09-302-491-5	Sequence 5, App1

539	27	58.7	498	2	US-09-202-491-6	Sequence 6, Appli	612	27	58.7	1437	2	US-09-647-140B-4	Sequence 4, Appli
540	27	58.7	498	2	US-08-817-318-2	Sequence 2, Appli	613	27	58.7	1437	2	US-09-528-031-2	Sequence 2, Appli
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752	26	56.5	221	2	US-09-248-796A-15088	Sequence 15088, A	825	26	56.5	336	2	US-10-134-297-8	Sequence 8, Appl1
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757	26	56.5	239	2	US-10-134-624-4	Sequence 4, Appl1	830	26	56.5	336	2	US-09-878-781-8	Sequence 8, Appl1

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990 26 56.5 485 2 US-09-009-494-2 Sequence 2, Appl
991 26 56.5 485 2 US-09-010-233-8 Sequence 8, Appl
992 26 56.5 485 2 US-09-010-232-4 Sequence 4, Appl
993 26 56.5 485 2 US-10-138-701-28 Sequence 28, Appl
994 26 56.5 490 2 US-09-134-000C-6176 Sequence 6176, Ap
995 26 56.5 491 2 US-09-248-796A-19609 Sequence 19609, A
996 26 56.5 493 2 US-09-252-991A-30722 Sequence 30722, A
997 26 56.5 493 2 US-09-489-039A-12903 Sequence 12903, A
998 26 56.5 487 2 US-09-949-016-69585 Sequence 6958, Ap
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ALIGNMENTS

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RESULT 1
US-07-909-122-3
; Sequence 3, Application US/07909122
; Patent No. 5415995
; GENERAL INFORMATION:
; APPLICANT: SCHOOLNIK, GARY K.
; APPLICANT: PALEFSKY, JOEL M.
; TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
; TITLE OF INVENTION: VIRUS
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 Page Mill Road
; CITY: Palo Alto
; STATE: California
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/909,122
; FILING DATE: 19920706
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: BENZ, WILLIAM H.
; REGISTRATION NUMBER: 25,952
; REFERENCE/DOCKET NUMBER: 28600-20105.01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 813-5600
; TELEFAX: (415) 494-0792
; TELEX: 706141
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 14 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-07-909-122-3
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Best Local Similarity 100.0%; Pred. No. 0.017;
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Db 2 ELQTTIHDI 10
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; Sequence 4, Application US/08363586
; Patent No. 5628161
; GENERAL INFORMATION:
; APPLICANT: Mueller, Martin
; APPLICANT: Giesmann, Lutz
; TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
; TITLE OF INVENTION: Peptides for the Diagnostic Purpose
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; STREET: 1300 I Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
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; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/363,586
; FILING DATE: 23-DEC-1994
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/909,296
; FILING DATE: 09-JUL-1992
; APPLICATION NUMBER: EP 91111720.8
; FILING DATE: 13-JUL-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Wadler, Linda A.
; REGISTRATION NUMBER: 33,218
; REFERENCE/DOCKET NUMBER: 02481-1195-00000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-408-4000
; TELEFAX: 202-408-4400
; INFORMATION FOR SEQ ID NO: 4:
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; TYPE: amino acid
; TOPOLOGY: linear
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; US-08-363-586-4
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; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCESCA
; APPLICANT: FERRIES, ESTELLE
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; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: WO/1 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-4

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; APPLICANT: INSTITUT OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
; FILE REFERENCE: N73477C GCW
; CURRENT APPLICATION NUMBER: US/09/701,080C
; CURRENT FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
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; LENGTH: 151
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; ORGANISM: Human papillomavirus
US-09-701-080C-18

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; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: WO/1 AO INS

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; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
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; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESS: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
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; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
US-08-316-239B-3

```

```

Query Match          100.0%; Score 46; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 1 ELQTTIHDI 9

```

Db 25 ELQTTIHDI 33

```
RESULT 7
US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Corsette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSER: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
; US-08-316-239B-4

Query Match 100.0%; Score 46; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 ELQTTIHDI 9
Db 25 ELQTTIHDI 33

RESULT 8
US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
```

```
; EARLIER APPLICATION NUMBER: AU PNO157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14
```

Query Match 100.0%; Score 46; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.25;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHDI 9
Db 94 ELQTTIHDI 102

```
RESULT 9
US-09-359-382-14
; Sequence 14, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-359-382-14
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Query Match 100.0%; Score 46; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.25;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHDI 9
Db 94 ELQTTIHDI 102

```
RESULT 10
US-08-117-083-10
; Sequence 10, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Bourneil, Michael E.
; APPLICANT: Ingilis, Stephen C.
; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; TITLE OF INVENTION: Papilloma Virus Proteins
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESSER: Walter H. Dreger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
```

COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,083
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-58783
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 182 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..182
OTHER INFORMATION: /note= "Xaa refers to stop codon in the open reading frame."
US-08-117-083-10

Query Match 100.0%; Score 46; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.26; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ELQTTIHDI 9
Db 26 ELQTTIHDI 34

RESULT 11
US-09-462-993-1
Sequence 1, Application US/09462993
Patent No. 6884786
GENERAL INFORMATION:
APPLICANT: KIENY, Marie-Paule
APPLICANT: BALLOU, Jean-Marc
APPLICANT: BIZOUARN, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
POLYPEPTIDE WITH MODIFIED CELL LOCATION
FILE REFERENCE: 017753-122
CURRENT APPLICATION NUMBER: US/09/462,993
CURRENT FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: PCT/FR98/01576
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
US-09-462-993-1

Query Match 100.0%; Score 46; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.36; Indels 0; Gaps 0;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ELQTTIHDI 9
Db 53 ELQTTIHDI 61

RESULT 12
US-08-860-165-10
Sequence 10, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PN0157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10

Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.39; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
Qy 1 ELQTTIHDI 9
Db 25 ELQTTIHDI 33

RESULT 13
US-09-359-382-10
Sequence 10, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PN0157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.39; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
Db 25 ELQTTTHDI 33

RESULT 14
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Patent No. 6428807
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; CURRENT FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 46; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.39;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
Db 25 ELQTTTHDI 33

RESULT 15
US-09-485-885-4
; Sequence 4, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 46; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
Db 131 ELQTTTHDI 139

RESULT 16
US-09-485-885-10
; Sequence 10, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 46; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.43;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
Db 150 ELQTTTHDI 158

RESULT 17
US-09-485-885-6
; Sequence 6, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; CURRENT FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 46; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
Db 131 ELQTTTHDI 139

RESULT 18
US-09-485-885-14
; Sequence 14, Application US/09485885

Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisee, Anne-Marie Eva Bernande
APPLICANT: Gerard, Catherine Marie Gristaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP96/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 46; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.59;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
DB 150 ELQTTIHDI 158

RESULT 19
US-08-787-547-102
Sequence 102, Application US/08787547
Patent No. 5783567
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Curley, Joanne M.
APPLICANT: Langer, Robert S.
TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY
NUMBER OF SEQUENCES: 107
CORRESPONDENCE ADDRESS:
ADDRESSER: Fish & Richardson, P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/787,547
FILING DATE: 22-JAN-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Frazer, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08191/003001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 102:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids

TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-787-547-102

Query Match 89.1%; Score 41; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
DB 1 LQTTIHDI 8

RESULT 20
US-08-159-339A-248
Sequence 248, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esben
TITLE OF INVENTION: HLA Binding peptides and Their
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 248:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-248

Query Match 89.1%; Score 41; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
DB 1 LQTTIHDI 8

RESULT 21
US-08-934-915-159
; Sequence 159, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USERFUL IN IMMUNOASSAY FOR
; NUMBER OF SEQUENCES: 193
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 1757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft Word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. Foutch
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946.6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:
; INFORMATION FOR SEQ ID NO: 159:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-934-915-159
Query Match 89.1%; Score 41; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 0.22;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
QY 1 ELQTTTHDI 9
Db 9 ELQTTTHNI 17
RESULT 22
US-08-913-159-14
; Sequence 14, Application US/08913159
; Patent No. 6300109
; GENERAL INFORMATION:
; APPLICANT:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USERFUL IN IMMUNOASSAY FOR
; NUMBER OF SEQUENCES: 14
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30 (EPO)

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/913,159
; FILING DATE:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: DK 0179/95
; FILING DATE: 17-FEB-1995
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 317 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-913-159-14
Query Match 78.3%; Score 36; DB 2; Length 317;
Best Local Similarity 75.0%; Pred. No. 37;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
QY 2 LQTTTHDI 9
Db 182 LKTTTHDV 189
RESULT 23
US-09-600-932-28
; Sequence 28, Application US/09600932
; Patent No. 6787639
; GENERAL INFORMATION:
; APPLICANT: Wakamiya, No. 6787639utaka
; TITLE OF INVENTION: NOVEL COLLECTIN
; FILE REFERENCE: 19036/36615
; CURRENT APPLICATION NUMBER: US/09/600,932
; CURRENT FILING DATE: 2000-09-08
; PRIOR APPLICATION NUMBER: PCT/JP98/03328
; PRIOR FILING DATE: 1998-07-24
; PRIOR APPLICATION NUMBER: JP 10-11281
; PRIOR FILING DATE: 1998-01-23
; NUMBER OF SEQ ID NOS: 29
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 28
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: surfactant protein A (SP-A)
US-09-600-932-28
Query Match 76.1%; Score 35; DB 2; Length 248;
Best Local Similarity 75.0%; Pred. No. 44;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
QY 1 ELQTTTHDI 8
Db 106 ELQATLHD 113
RESULT 24
US-09-949-016-6612
; Sequence 6612, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CL001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08

; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6612
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-6612

Query Match 76.1%; Score 35; DB 2; Length 248;
Best Local Similarity 75.0%; Pred. No. 44;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
|||:|
Db 106 ELQATLHD 113

RESULT 25
US-09-949-016-8481
; Sequence 8481, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: C1001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; PRIOR FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8481
; LENGTH: 251
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-8481

Query Match 76.1%; Score 35; DB 2; Length 251;
Best Local Similarity 75.0%; Pred. No. 45;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
|||:|
Db 109 ELQATLHD 116

RESULT 26
US-09-949-016-9058
; Sequence 9058, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: C1001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; PRIOR FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 9058
; LENGTH: 251
; TYPE: PRT

; ORGANISM: Human
US-09-949-016-9058

Query Match 76.1%; Score 35; DB 2; Length 251;
Best Local Similarity 75.0%; Pred. No. 45;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
|||:|
Db 109 ELQATLHD 116

RESULT 27
US-09-949-016-9059
; Sequence 9059, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: C1001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; PRIOR FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 9059
; LENGTH: 259
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-9059

Query Match 76.1%; Score 35; DB 2; Length 259;
Best Local Similarity 75.0%; Pred. No. 46;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
|||:|
Db 117 ELQATLHD 124

RESULT 28
US-08-936-165A-297
; Sequence 297, Application US/08936165A
; Patent No. 6348582
; GENERAL INFORMATION:
; APPLICANT: Black, Michael
; APPLICANT: Burnham, Martin
; APPLICANT: Hodgson, John
; APPLICANT: Knowles, David
; APPLICANT: Lometto, Michael
; APPLICANT: Nicholas, Richard
; APPLICANT: Pratt, Julie
; APPLICANT: Reichard, Richard
; APPLICANT: Rosenberg, Martin
; APPLICANT: Ward, Judith
; TITLE OF INVENTION: No. 6348582el Prokaryotic polynucleotides,
; TITLE OF INVENTION: Polypeptides and their uses
; NUMBER OF SEQUENCES: 534
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: SmithKline Beecham Corporation
; STREET: 709 Swedeland Road
; CITY: King of Prussia
; STATE: PA
; COUNTRY: USA
; ZIP: 19406-0939
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette

```
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/936.165A
FILING DATE: 24-SEP-1997
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/027,032
FILING DATE: 24-SEP-1996
ATTORNEY/AGENT INFORMATION:
NAME: Gimmil, Edward R
REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: P50549
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
TELEFAX: 610-270-5090
TELEX:
INFORMATION FOR SEQ ID NO: 297:
SEQUENCE CHARACTERISTICS:
LENGTH: 56 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Protein
US-08-936-165A-297
```

```
Query Match      73.9%; Score 34; DB 2; Length 56;
Best Local Similarity 66.7%; Pred. No. 14;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTTIHD 9
DB      6 EMKRTIHD 14
```

```
RESULT 29
US-09-603-208A-62
Sequence 62, Application US/09603208A
Patent No. 6822084
GENERAL INFORMATION:
APPLICANT: Pompeju, Markus
APPLICANT: Krogger, Burhard
APPLICANT: Schroder, Hartwig
APPLICANT: Zelder, Oskar
APPLICANT: Habernauer, Gregor
APPLICANT: Lee, Heung-Shick
TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING STRESS,
FILE REFERENCE: BGI-124CP
CURRENT APPLICATION NUMBER: US/09/603,208A
CURRENT FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: 60/141031
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 60/142692
PRIOR FILING DATE: 1998-07-01
PRIOR APPLICATION NUMBER: 60/151214
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: DE 19930429.7
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: DE 19931413.6
PRIOR FILING DATE: 1998-07-08
PRIOR APPLICATION NUMBER: DE 19931457.8
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19931541.8
PRIOR FILING DATE: 1998-07-08
PRIOR APPLICATION NUMBER: DE 19932209.0
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: DE 19932230.9
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: DE 19932914.1
PRIOR FILING DATE: 1999-07-14
```

```
PRIOR APPLICATION NUMBER: DE 19940764.9
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: DE 19941382.7
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 306
SEQ ID NO 62
LENGTH: 383
TYPE: PRT
ORGANISM: Corynebacterium glutamicum
US-09-603-208A-62
```

```
Query Match      73.9%; Score 34; DB 2; Length 383;
Best Local Similarity 66.7%; Pred. No. 1.1e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTTIHD 9
DB      237 EMKRTIHD 245
```

```
RESULT 30
US-09-949-016-7565
Sequence 7565, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 7565
LENGTH: 640
TYPE: PRT
ORGANISM: Human
US-09-949-016-7565
```

```
Query Match      73.9%; Score 34; DB 2; Length 640;
Best Local Similarity 75.0%; Pred. No. 1.9e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTTIHD 8
DB      99 ELQVTMHD 106
```

```
RESULT 31
US-09-248-796A-20895
Sequence 20895, Application US/09248796A
Patent No. 6747137
GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICA
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 20895
LENGTH: 695
TYPE: PRT
```


ORGANISM: Candida albicans
US-09-248-796A-20895

Query Match 73.9%; Score 34; DB 2; Length 695;
Best Local Similarity 66.7%; Pred. No. 2.1e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
|:|:|:|:
DB 28 EVQTTVEDI 36

RESULT 32
US-09-252-991A-22216
Sequence 22216, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIOR APPLICATION NUMBER: US 60/074,788
PRIOR FILING DATE: 1998-02-18
PRIOR APPLICATION NUMBER: US 60/094,190
PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 31142
SEQ ID NO 22216
LENGTH: 811
TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-22216

Query Match 73.9%; Score 34; DB 2; Length 811;
Best Local Similarity 75.0%; Pred. No. 2.4e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 8
|:|:|:|:
DB 792 ELQTTLHD 799

RESULT 33
US-09-964-956-13
Sequence 13, Application US/09964956
Patent No. 6875570
GENERAL INFORMATION:
APPLICANT: Gerlach, Valerie L
APPLICANT: Macdougall, John R
APPLICANT: Smithson, Glenda
APPLICANT: Miller, Isabelle
APPLICANT: Stone, David
APPLICANT: Gunther, Erik
APPLICANT: Ellerman, Karen
APPLICANT: Grose, William M
APPLICANT: Alsobrook II, John P
APPLICANT: Lepley, Denise M
APPLICANT: Burgess, Catherine B
APPLICANT: Padigaru, Muralidhara
APPLICANT: Kekuda, Ramesh
APPLICANT: Spytek, Kimberly A
APPLICANT: Leach, Martin D
APPLICANT: Shimkets, Richard A
TITLE OF INVENTION: No. 6875570e1 Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-124
CURRENT APPLICATION NUMBER: US/09/964,956
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/235,631
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/235,633
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/235,808

PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,064
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,065
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,066
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,135
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: 60/237,434
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/238,321
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/238,399
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 60/238,396
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 60/276,667
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/294,823
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/304,868
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 127
SOFTWARE: Patentln Ver. 2.1
SEQ ID NO 13
LENGTH: 1896
TYPE: PRT
ORGANISM: Homo sapiens
US-09-964-956-13

Query Match 73.9%; Score 34; DB 2; Length 1896;
Best Local Similarity 66.7%; Pred. No. 6e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
|:|:|:|:
DB 1295 ELQTTDHEL 1303

RESULT 34
US-09-964-956-44
Sequence 44, Application US/09964956
Patent No. 6875570
GENERAL INFORMATION:
APPLICANT: Gerlach, Valerie L
APPLICANT: Macdougall, John R
APPLICANT: Smithson, Glenda
APPLICANT: Miller, Isabelle
APPLICANT: Stone, David
APPLICANT: Gunther, Erik
APPLICANT: Ellerman, Karen
APPLICANT: Grose, William M
APPLICANT: Alsobrook II, John P
APPLICANT: Lepley, Denise M
APPLICANT: Burgess, Catherine B
APPLICANT: Padigaru, Muralidhara
APPLICANT: Kekuda, Ramesh
APPLICANT: Spytek, Kimberly A
APPLICANT: Leach, Martin D
APPLICANT: Shimkets, Richard A
TITLE OF INVENTION: No. 6875570e1 Proteins and Nucleic Acids Encoding Same
FILE REFERENCE: 21402-124
CURRENT APPLICATION NUMBER: US/09/964,956
CURRENT FILING DATE: 2001-09-26
PRIOR APPLICATION NUMBER: 60/235,631
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/235,633
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/235,808
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,064

PRIOR APPLICATION NUMBER: 60/236,065
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,066
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: 60/236,135
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: 60/237,434
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/238,321
PRIOR FILING DATE: 2000-10-05
PRIOR APPLICATION NUMBER: 60/238,399
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 60/238,396
PRIOR FILING DATE: 2000-10-06
PRIOR APPLICATION NUMBER: 60/276,667
PRIOR FILING DATE: 2001-03-16
PRIOR APPLICATION NUMBER: 60/294,823
PRIOR FILING DATE: 2001-05-31
PRIOR APPLICATION NUMBER: 60/304,868
PRIOR FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 127
SOFTWARE: Patent Ver. 2.1
SEQ ID NO: 44
LENGTH: 1905
TYPE: PRT
ORGANISM: Xenopus laevis
US-09-964-956-44

Query Match 73.9%; Score 34; DB 2; Length 1905;
Best Local Similarity 66.7%; Pred. No. 6e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 1292 ELQTTIHDL 1300

RESULT 35
US-08-118-270-256
Sequence 256, Application US/08118270
Patent No. 5508384
GENERAL INFORMATION:
APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
City: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY-2A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
TELEX: 248633

INFORMATION FOR SEQ ID NO: 256:
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-118-270-256

Query Match 71.7%; Score 33; DB 1; Length 59;
Best Local Similarity 55.6%; Pred. No. 23;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 25 ELSTIHDL 33

RESULT 36
PCT-US93-08528-256
Sequence 256, Application PC/TUS9308528
GENERAL INFORMATION:
APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
City: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08528
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY-2 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO: 256:
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
PCT-US93-08528-256

Query Match 71.7%; Score 33; DB 4; Length 59;
Best Local Similarity 55.6%; Pred. No. 23;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 25 ELSTIHDL 33

RESULT 37
US-09-270-767-33046
Sequence 33046, Application US/09270767

Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 33046
; LENGTH: 166
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-33046

Query Match 71.7%; Score 33; DB 2; Length 166;
Best Local Similarity 62.5%; Pred. No. 69;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
|:|:|:|:
Db 85 LRTTIDHL 92

RESULT 38
US-09-270-767-48263
; Sequence 48263, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 48263
; LENGTH: 166
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-48263

Query Match 71.7%; Score 33; DB 2; Length 166;
Best Local Similarity 62.5%; Pred. No. 69;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
|:|:|:|:
Db 85 LRTTIDHL 92

RESULT 39
US-09-134-001C-4050
; Sequence 4050, Application US/09134001C
; Patent No. 6380370
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCUS
; FILE REFERENCE: GTC-007
; CURRENT APPLICATION NUMBER: US/09/134,001C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/064,964
; PRIOR FILING DATE: 1997-11-08
; PRIOR APPLICATION NUMBER: US 60/055,779
; PRIOR FILING DATE: 1997-08-14
; NUMBER OF SEQ ID NOS: 5674
; SEQ ID NO 4050
; LENGTH: 198
; TYPE: PRT
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-4050

Query Match 71.7%; Score 33; DB 2; Length 198;
Best Local Similarity 62.5%; Pred. No. 83;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
|:|:|:|:
Db 128 EINTTIDHD 135

RESULT 40
US-09-354-123-6
; Sequence 6, Application US/09354123
; Patent No. 6489537
; GENERAL INFORMATION:
; APPLICANT: Rea, Philip A.
; APPLICANT: Vatamanluk, Olena K.
; APPLICANT: Mari, Stephanie
; APPLICANT: Lu, Yu-Ping
; APPLICANT: Schroeder, Julian I.
; APPLICANT: Kim, Eugene J.
; APPLICANT: Clemens, Stephan
; TITLE OF INVENTION: NOVEL PHYTOCHELATIN SYNTHASES AND USES THEREFOR
; FILE REFERENCE: 9596-102U1/209596,0289
; CURRENT APPLICATION NUMBER: US/09/354,123
; CURRENT FILING DATE: 1999-07-15
; EARLIER APPLICATION NUMBER: 09/315,449
; EARLIER FILING DATE: 1999-05-20
; EARLIER APPLICATION NUMBER: 60/095,624
; EARLIER FILING DATE: 1998-08-07
; NUMBER OF SEQ ID NOS: 10
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 6
; LENGTH: 500
; TYPE: PRT
; ORGANISM: Triticum aestivum
US-09-354-123-6

Query Match 71.7%; Score 33; DB 2; Length 500;
Best Local Similarity 100.0%; Pred. No. 2.2e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 QTTIHD 8
|:|:|:|:
Db 125 QTTIHD 130

RESULT 41
US-10-214-269-19
; Sequence 19, Application US/10214269
; Patent No. 6844485
; GENERAL INFORMATION:
; APPLICANT: Butler, Karlene H.
; APPLICANT: Famodu, Omolayo O.
; APPLICANT: Harwell, Leslie T.
; APPLICANT: Orozco, Jr., Emil M.
; APPLICANT: Rasco-Gaunt, Soniriza
; APPLICANT: Thorpe, Catherine J.
; TITLE OF INVENTION: phytochelatn synthase
; FILE REFERENCE: BB1511 US NA
; CURRENT APPLICATION NUMBER: US/10/214,269
; CURRENT FILING DATE: 2002-08-07
; PRIOR APPLICATION NUMBER: US 60/310,521
; PRIOR FILING DATE: 2001-08-07
; NUMBER OF SEQ ID NOS: 21
; SOFTWARE: Microsoft Office 97
; SEQ ID NO 19
; LENGTH: 500
; TYPE: PRT
; ORGANISM: Triticum aestivum
US-10-214-269-19

Query Match 71.7%; Score 33; DB 2; Length 500;
Best Local Similarity 100.0%; Pred. No. 2.2e+02;

Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 QTTIHD 8
Db 125 QTTIHD 130

RESULT 42
US-09-990-337-3

; Sequence 3, Application US/09990337
; Patent No. 6703223
; GENERAL INFORMATION:
; APPLICANT: BATHIE, Brigitte
; APPLICANT: SCHISCHKA, Natalie
; APPLICANT: BOTT, Michael
; APPLICANT: SCHAEFER, Steffen
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES CODING FOR THE MIRA and/or MxRb PROTEINS
; FILE REFERENCE: 215474US0X
; CURRENT APPLICATION NUMBER: US/09/990,337
; PRIOR FILING DATE: 2001-11-23
; PRIOR APPLICATION NUMBER: DE10057802.0
; PRIOR FILING DATE: 2000-11-22
; PRIOR APPLICATION NUMBER: DE10125089.4
; PRIOR FILING DATE: 2001-05-23
; NUMBER OF SEQ ID NOS: 7
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 503
; TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
US-09-990-337-3

Query Match
Best Local Similarity 71.7%; Score 33; DB 2; Length 503;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 ELQTTIHD 9
Db 326 ELSTALHDV 334

RESULT 43
US-09-543-681A-7528
; Sequence 7528, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; PRIOR FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO 7528
; LENGTH: 738
; TYPE: PRT
; ORGANISM: Proteus mirabilis
US-09-543-681A-7528

Query Match
Best Local Similarity 71.7%; Score 33; DB 2; Length 738;
Matches 7; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 ELQTTIHD 9
Db 606 ELQTTEDI 614

RESULT 44
US-09-248-796A-20513
; Sequence 20513, Application US/09248796A

; Patent No. 6747137
; GENERAL INFORMATION:

; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICA
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; PRIOR FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 20513
; LENGTH: 933
; TYPE: PRT
; ORGANISM: Candida albicans
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (635)
; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unknown
US-09-248-796A-20513

Query Match
Best Local Similarity 71.7%; Score 33; DB 2; Length 933;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 9
Db 263 EVATTLHDL 271

RESULT 45
US-09-252-991A-31032
; Sequence 31032, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 31032
; LENGTH: 1128
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-31032

Query Match
Best Local Similarity 71.7%; Score 33; DB 2; Length 1128;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
Db 1010 ELRTTVHE 1017

RESULT 46
US-09-949-016-8280
; Sequence 8280, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: CLO01307
; CURRENT APPLICATION NUMBER: US/09/949,016

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; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8280
; LENGTH: 1769
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-8280
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Query Match          71.7%; Score 33; DB 2; Length 1769;
Best Local Similarity 55.6%; Pred. No. 8.6e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;
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```
Qy      1 ELQTTIHDI 9
      ||:|:|:|:|:
Db      385 ELRTIVHDL 393
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```
RESULT 47
US-09-949-016-8281
; Sequence 8281, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: CLO01307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8281
; LENGTH: 1769
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-8281
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```
Query Match          71.7%; Score 33; DB 2; Length 1769;
Best Local Similarity 55.6%; Pred. No. 8.6e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;
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```
Qy      1 ELQTTIHDI 9
      ||:|:|:|:|:
Db      385 ELRTIVHDL 393
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```
RESULT 48
US-09-949-016-8282
; Sequence 8282, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; FILE REFERENCE: CLO01307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
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; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 8282
; LENGTH: 1769
; TYPE: PRT
; ORGANISM: Human
US-09-949-016-8282
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Query Match          71.7%; Score 33; DB 2; Length 1769;
Best Local Similarity 55.6%; Pred. No. 8.6e+02;
Matches      5; Conservative      3; Mismatches      1; Indels      0; Gaps      0;
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```
Qy      1 ELQTTIHDI 9
      ||:|:|:|:|:
Db      385 ELRTIVHDL 393
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RESULT 49
US-09-040-738-2
; Sequence 2, Application US/09040738
; Patent No. 6207374
; GENERAL INFORMATION:
; APPLICANT: Sampson et al.
; TITLE OF INVENTION: Tubercous Sclerosis 2 Gene and Uses
; TITLE OF INVENTION: Thereof
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Kathleen M. Williams, Banner & Witcoff,
; STREET: One Financial Center
; CITY: Boston
; STATE: Massachusetts
; COUNTRY: USA
; ZIP: 02111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Wordperfect 6.1a
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/09/040,738
; FILING DATE: Concurrently herewith
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: GB9326470.3
; FILING DATE: 24-December-1993
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: GB9411900.5
; FILING DATE: 14-June-1994
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/GB94/02823
; FILING DATE: 23-December-1994
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/652,426
; FILING DATE: 30-May-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Williams, Kathleen M.
; REGISTRATION NUMBER: 34,380
; REFERENCE/DOCKET NUMBER: 3265/73963
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-345-9100
; TELEFAX: 617-345-9111
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1784 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-09-040-738-2
```

```
Query Match          71.7%; Score 33; DB 2; Length 1784;
```

Best Local Similarity 55.6%; Pred. No. 8.7e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD1 9
||:|:|:
Db 379 ELRTIVHDL 387

RESULT 50
US-08-652-426A-2
; Sequence 2, Application US/08652426A
; Patent No. 6232452
; GENERAL INFORMATION:
; APPLICANT: Sampson et al.
; TITLE OF INVENTION: Tuberos Sclerosis 2 Gene and Uses
; TITLE OF INVENTION: Thereof
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Kathleen M. Williams, Banner & Wilcoff,
; STREET: One Financial Center
; CITY: Boston
; STATE: Massachusetts
; COUNTRY: USA
; ZIP: 02111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Wordperfect 6.1a
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/652,426A
; FILING DATE: 01-October-1996
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: GB9326470.3
; FILING DATE: 12/24/93
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: GB9411900.5
; FILING DATE: 06/14/94
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: PCT/GB94/02823
; FILING DATE: 12/23/94
; ATTORNEY/AGENT INFORMATION:
; NAME: Williams, Kathleen M.
; REGISTRATION NUMBER: 34,380
; REFERENCE/DOCKET NUMBER: 96,738
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-345-9100
; TELEFAX: 617-345-9111
; INFORMATION FOR SEQ ID NO: 2:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 1784 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-652-426A-2

Query Match 71.7%; Score 33; DB 2; Length 1784;
Best Local Similarity 55.6%; Pred. No. 8.7e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
QY 1 ELQTTIHD1 9
||:|:|:
Db 379 ELRTIVHDL 387

Search completed: May 5, 2006, 03:13:19
Job time : 22.7 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 07:56:48 ; Search time 56 Seconds
(without alignments)
67.151 Million cell updates/sec

Title: US-08-170-344-5
Perfect score: 46
Sequence: 1 ELQTTIHDI 9

Scoring table: BIOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :
1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep.*
2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
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2	46	100.0	15	US-10-476-570-21	Sequence 21, Appli
3	46	100.0	15	US-10-476-570-22	Sequence 22, Appli
4	46	100.0	21	US-10-476-570-8	Sequence 8, Appli
5	46	100.0	25	US-11-021-949-1	Sequence 53, Appli
6	46	100.0	30	US-10-476-570-53	Sequence 4, Appli
7	46	100.0	30	US-10-858-384-4	Sequence 9, Appli
8	46	100.0	32	US-10-476-570-19	Sequence 19, Appli
9	46	100.0	33	US-10-476-570-9	Sequence 6, Appli
10	46	100.0	151	US-10-484-063-20	Sequence 20, Appli
11	46	100.0	151	US-10-484-063-27	Sequence 27, Appli
12	46	100.0	151	US-10-858-384-2	Sequence 2, Appli
13	46	100.0	158	US-10-367-057-16	Sequence 16, Appli
14	46	100.0	158	US-11-021-949-13	Sequence 13, Appli
15	46	100.0	158	US-10-472-724-2	Sequence 2, Appli
16	46	100.0	171	US-11-072-288-1	Sequence 1, Appli
17	46	100.0	243	US-09-367-309A-1	Sequence 1, Appli
18	46	100.0	266	US-10-000-903-4	Sequence 4, Appli
19	46	100.0	273	US-10-899-771-4	Sequence 4, Appli
20	46	100.0	273	US-10-899-771-10	Sequence 10, Appli
21	46	100.0	292	US-10-899-771-10	Sequence 10, Appli
22	46	100.0	371	US-10-899-771-6	Sequence 6, Appli
23	46	100.0	371	US-10-899-771-6	Sequence 6, Appli
24	46	100.0	390	US-10-000-903-14	Sequence 14, Appli
25	46	100.0	390	US-10-899-771-14	Sequence 14, Appli
26	46	100.0	536	US-10-367-095-10	Sequence 10, Appli
27	46	100.0	536	US-10-367-095-10	Sequence 10, Appli

28	46	100.0	536	US-10-368-046-10	Sequence 10, Appli
29	46	100.0	536	US-10-367-367-10	Sequence 10, Appli
30	46	100.0	536	US-10-318-337-10	Sequence 10, Appli
31	42	91.3	248	US-10-820-155-42	Sequence 42, Appli
32	42	91.3	248	US-10-820-155-57	Sequence 57, Appli
33	32	41	9	US-09-909-460-102	Sequence 102, App
34	34	41	9	US-09-872-836-102	Sequence 102, App
35	41	89.1	9	US-10-133-210-279	Sequence 279, App
36	41	89.1	9	US-10-758-870-102	Sequence 102, App
37	41	89.1	9	US-10-751-845-56	Sequence 56, Appli
38	37	80.4	1198	US-10-452-024-95	Sequence 95, Appli
39	36	78.3	151	US-10-476-570-20	Sequence 20, Appli
40	36	78.3	151	US-10-425-115-237192	Sequence 237192, A
41	36	78.3	154	US-11-097-143-21504	Sequence 21504, A
42	35	76.1	93	US-10-312-829-6	Sequence 6, Appli
43	35	76.1	145	US-09-925-302-845	Sequence 845, App
44	35	76.1	145	US-09-925-302-845	Sequence 845, App
45	35	76.1	148	US-10-312-829-5	Sequence 5, Appli
46	35	76.1	166	US-10-336-603A-98	Sequence 98, Appli
47	35	76.1	223	US-10-437-963-163049	Sequence 163049, A
48	35	76.1	243	US-10-336-603A-96	Sequence 96, Appli
49	35	76.1	248	US-10-312-829-4	Sequence 4, Appli
50	35	76.1	248	US-10-336-603A-100	Sequence 100, App
51	35	76.1	248	US-10-971-461-25	Sequence 25, Appli
52	35	76.1	248	US-10-971-461-25	Sequence 26, Appli
53	35	76.1	248	US-10-820-155-58	Sequence 58, Appli
54	35	76.1	248	US-10-756-149-5337	Sequence 5337, App
55	35	76.1	259	US-09-925-302-473	Sequence 473, App
56	35	76.1	259	US-09-925-302-473	Sequence 297, App
57	34	73.9	56	US-09-939-980-297	Sequence 50864, A
58	34	73.9	97	US-10-767-971-50864	Sequence 335035, A
59	34	73.9	122	US-10-425-115-335035	Sequence 443, App
60	34	73.9	129	US-09-925-298-443	Sequence 443, App
61	34	73.9	129	US-10-102-806-443	Sequence 263392, A
62	34	73.9	156	US-10-425-115-263392	Sequence 361, App
63	34	73.9	158	US-11-021-949-161	Sequence 293053, A
64	34	73.9	159	US-10-425-115-293053	Sequence 60280, A
65	34	73.9	166	US-10-425-114-60280	Sequence 342845, A
66	34	73.9	234	US-10-425-115-342845	Sequence 191647, A
67	34	73.9	248	US-10-424-599-191647	Sequence 71160, A
68	34	73.9	259	US-10-425-114-71760	Sequence 52268, A
69	34	73.9	309	US-10-450-763-52268	Sequence 168462, A
70	34	73.9	316	US-10-424-599-168462	Sequence 117225, A
71	34	73.9	403	US-10-437-963-117225	Sequence 263390, A
72	34	73.9	448	US-10-425-115-263390	Sequence 49312, A
73	34	73.9	463	US-10-425-114-49312	Sequence 9, Appli
74	34	73.9	463	US-10-844-716-9	Sequence 2369, App
75	34	73.9	595	US-10-276-774-3269	Sequence 3259, App
76	34	73.9	619	US-10-320-797-3259	Sequence 9175, App
77	34	73.9	623	US-10-739-930-9175	Sequence 106, App
78	34	73.9	628	US-10-176-847-108	Sequence 246, App
79	34	73.9	628	US-10-205-823-246	Sequence 6, Appli
80	34	73.9	628	US-10-257-021-6	Sequence 70, Appli
81	34	73.9	628	US-10-257-021-6	Sequence 52, Appli
82	34	73.9	628	US-10-789-378-52	Sequence 246, App
83	34	73.9	628	US-11-051-454-246	Sequence 49664, A
84	34	73.9	1082	US-10-282-122A-49664	Sequence 82, Appli
85	34	73.9	1754	US-10-087-664-82	Sequence 81, Appli
86	34	73.9	1754	US-10-218-779-82	Sequence 81, Appli
87	34	73.9	1894	US-10-087-664-81	Sequence 81, Appli
88	34	73.9	1894	US-10-218-779-81	Sequence 9, Appli
89	34	73.9	1894	US-10-451-010-9	Sequence 13, Appli
90	34	73.9	1896	US-09-964-956-13	Sequence 34, Appli
91	34	73.9	1896	US-10-312-352-34	Sequence 44, Appli
92	34	73.9	1905	US-09-964-956-44	Sequence 32, Appli
93	34	73.9	1905	US-10-087-664-32	Sequence 204, App
94	34	73.9	1925	US-10-218-779-32	Sequence 315095, A
95	33	71.7	22	US-10-639-067-204	Sequence 368030, A
96	33	71.7	40	US-10-425-115-315095	Sequence 16, Appli
97	33	71.7	102	US-10-425-115-368030	Sequence 360, App
98	33	71.7	149	US-11-021-949-16	Sequence 31, Appli
99	33	71.7	149	US-11-021-949-360	
100	33	71.7	162	US-11-021-949-31	

101	33	71.7	198	4	US-10-724-972A-4917	Sequence 4917, Ap	174	32	69.6	1104	4	US-10-369-493-22548	Sequence 22548, A
102	33	71.7	233	4	US-10-772-272A-16	Sequence 16, Appl	175	32	69.6	1258	4	US-10-310-154-703	Sequence 703, App
103	33	71.7	247	5	US-10-820-155-44	Sequence 44, Appl	176	32	69.6	1258	5	US-10-732-923-621	Sequence 621, App
104	33	71.7	247	5	US-10-820-155-73	Sequence 73, Appl	177	32	69.6	1258	5	US-10-732-923-22596	Sequence 22596, A
105	33	71.7	247	5	US-10-820-155-78	Sequence 78, Appl	178	32	69.6	1280	6	US-11-047-143-1539	Sequence 1539, App
106	33	71.7	406	4	US-10-369-493-12152	Sequence 12152, A	179	32	69.6	1280	6	US-10-745-237-180	Sequence 180, App
107	33	71.7	499	3	US-09-738-626-3342	Sequence 4342, A	180	32	69.6	1351	6	US-11-097-143-36795	Sequence 36795, A
108	33	71.7	499	6	US-11-006-098-30	Sequence 30, Appl	181	31	67.4	46	4	US-10-424-599-230580	Sequence 230580, A
109	33	71.7	500	4	US-10-214-269-19	Sequence 19, Appl	182	31	67.4	48	4	US-10-424-599-157049	Sequence 157049, A
110	33	71.7	503	3	US-09-990-337-3	Sequence 3, Appl1	183	31	67.4	49	4	US-10-437-963-157793	Sequence 157793, A
111	33	71.7	503	4	US-10-411-318-3	Sequence 3, Appl1	184	31	67.4	48	4	US-10-424-599-258195	Sequence 258195, A
112	33	71.7	503	5	US-10-651-967-3	Sequence 3, Appl1	185	31	67.4	51	4	US-10-425-115-196748	Sequence 196748, A
113	33	71.7	568	6	US-11-097-143-28185	Sequence 28185, A	186	31	67.4	68	4	US-10-437-963-163168	Sequence 163168, A
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115	33	71.7	908	5	US-10-805-684-87	Sequence 87, Appl	188	31	67.4	78	4	US-10-424-599-284841	Sequence 284841, A
116	33	71.7	1130	5	US-10-260-708-57	Sequence 67, Appl	189	31	67.4	81	4	US-10-425-115-194108	Sequence 194108, A
117	33	71.7	1317	3	US-09-815-242-5118	Sequence 5118, Ap	190	31	67.4	87	4	US-10-424-599-209998	Sequence 209998, A
118	33	71.7	1317	4	US-10-282-122A-43495	Sequence 43495, A	191	31	67.4	112	4	US-10-767-701-38587	Sequence 38587, A
119	33	71.7	1585	4	US-10-437-963-124349	Sequence 124349, A	192	31	67.4	121	6	US-11-097-143-5853	Sequence 5853, Ap
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121	33	71.7	1882	5	US-10-450-763-35443	Sequence 35443, A	194	31	67.4	146	4	US-10-424-599-278042	Sequence 278042, A
122	33	71.7	1886	5	US-10-450-763-38020	Sequence 38020, A	195	31	67.4	157	4	US-10-424-599-27980	Sequence 27980, A
123	33	71.7	2265	4	US-10-092-900A-296	Sequence 296, App	196	31	67.4	165	4	US-10-425-114-55470	Sequence 55470, A
124	33	71.7	2296	5	US-10-723-860-1555	Sequence 1555, Ap	197	31	67.4	177	4	US-10-425-115-277091	Sequence 277091, A
125	33	71.7	2296	5	US-10-756-149-5112	Sequence 5112, Ap	198	31	67.4	183	4	US-10-029-386-32448	Sequence 32448, A
126	33	71.7	2327	4	US-10-092-900A-292	Sequence 294, App	199	31	67.4	197	4	US-10-282-122A-61172	Sequence 61172, A
127	33	71.7	2390	4	US-10-092-900A-292	Sequence 292, App	200	31	67.4	208	4	US-10-335-977-9731	Sequence 9731, Ap
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131	32	69.6	77	4	US-10-106-698-5327	Sequence 5327, Ap	204	31	67.4	332	4	US-10-369-493-8092	Sequence 8092, Ap
132	32	69.6	137	4	US-10-437-963-161775	Sequence 161775, A	205	31	67.4	336	4	US-10-424-599-236648	Sequence 236648, A
133	32	69.6	148	4	US-10-437-963-201751	Sequence 201751, A	206	31	67.4	345	4	US-10-369-493-6577	Sequence 6577, Ap
134	32	69.6	163	4	US-10-767-701-51443	Sequence 51443, A	207	31	67.4	345	4	US-10-369-493-6577	Sequence 6577, Ap
135	32	69.6	164	4	US-10-425-115-186341	Sequence 186341, A	208	31	67.4	359	4	US-10-332-512A-16	Sequence 16, Appl
136	32	69.6	164	4	US-10-425-115-186344	Sequence 186344, A	209	31	67.4	364	4	US-10-425-115-287375	Sequence 287375, A
137	32	69.6	164	4	US-10-425-115-186347	Sequence 186347, A	210	31	67.4	369	4	US-10-425-115-299865	Sequence 299865, A
138	32	69.6	166	4	US-10-437-963-172480	Sequence 172480, A	211	31	67.4	371	4	US-10-437-963-189109	Sequence 189109, A
139	32	69.6	212	4	US-10-437-963-145884	Sequence 145884, A	212	31	67.4	387	4	US-10-425-114-72757	Sequence 72757, A
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141	32	69.6	242	3	US-09-728-626-6004	Sequence 6004, Ap	214	31	67.4	395	4	US-10-841-707-8	Sequence 8, Appl1
142	32	69.6	246	4	US-10-424-599-168148	Sequence 168148, A	215	31	67.4	398	4	US-10-425-114-55344	Sequence 55344, A
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144	32	69.6	331	5	US-10-450-763-47010	Sequence 47010, A	217	31	67.4	417	4	US-10-424-599-186616	Sequence 186616, A
145	32	69.6	333	4	US-10-369-493-10175	Sequence 10175, A	218	31	67.4	417	4	US-10-424-599-271037	Sequence 271037, A
146	32	69.6	335	4	US-10-369-493-20899	Sequence 20899, A	219	31	67.4	418	3	US-09-927-602-5	Sequence 5, Appl1
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153	32	69.6	421	5	US-10-450-763-50008	Sequence 50008, A	226	31	67.4	433	2	US-08-945-038-6	Sequence 6, Appl1
154	32	69.6	469	5	US-10-732-923-6198	Sequence 6198, Ap	227	31	67.4	433	4	US-10-425-114-63366	Sequence 63366, A
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156	32	69.6	501	5	US-10-732-923-9631	Sequence 9631, Ap	229	31	67.4	434	4	US-10-437-963-149104	Sequence 149104, A
157	32	69.6	501	5	US-10-732-923-9692	Sequence 9692, Ap	230	31	67.4	435	4	US-10-425-114-59462	Sequence 59462, A
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160	32	69.6	513	5	US-10-472-928-2686	Sequence 2686, Ap	233	31	67.4	537	4	US-10-424-599-230650	Sequence 230650, A
161	32	69.6	531	4	US-10-352-839-7	Sequence 7, Appl1	234	31	67.4	540	4	US-10-424-599-230656	Sequence 230656, A
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163	32	69.6	553	4	US-10-467-534-61	Sequence 61, Appl	236	31	67.4	552	4	US-10-425-114-46347	Sequence 46347, A
164	32	69.6	558	4	US-10-205-194-154	Sequence 154, App	237	31	67.4	552	4	US-10-425-114-46829	Sequence 46829, A
165	32	69.6	588	5	US-10-450-763-51492	Sequence 51492, A	238	31	67.4	564	4	US-10-424-599-272057	Sequence 272057, A
166	32	69.6	578	5	US-10-617-038-41	Sequence 41, Appl	239	31	67.4	652	4	US-10-467-490-5	Sequence 5, Appl1
167	32	69.6	601	6	US-11-097-143-4221	Sequence 4221, App	240	31	67.4	785	5	US-10-264-04A-2968	Sequence 2968, Ap
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171	32	69.6	764	4	US-10-437-963-151490	Sequence 151490, A	244	31	67.4	841	4	US-10-467-490-2	Sequence 2, Appl1
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173	32	69.6	1052	5	US-10-450-763-35824	Sequence 35824, A	246	31	67.4	846	4	US-10-072-841-33	Sequence 33, Appl

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248	31	67.4	846	5	US-10-894-643-33	Sequence 33, Appl	321	30	65.2	342	4	US-10-425-114-6858	Sequence 6858, A
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256	31	67.4	995	4	US-10-094-749-2626	Sequence 2626, Ap	329	30	65.2	419	4	US-10-437-963-19308	Sequence 19308, A
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279	30	65.2	112	3	US-09-893-737-322	Sequence 322, Ap	352	30	65.2	456	5	US-10-719-993-671	Sequence 671, Ap
280	30	65.2	112	5	US-10-970-713-322	Sequence 322, Ap	353	30	65.2	456	5	US-11-097-143-21840	Sequence 21840, A
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285	30	65.2	123	4	US-10-425-115-503420	Sequence 503420, A	358	30	65.2	456	5	US-10-425-114-63473	Sequence 63473, A
286	30	65.2	126	4	US-10-767-701-50300	Sequence 50300, A	359	30	65.2	456	5	US-10-369-493-6662	Sequence 6662, Ap
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297	30	65.2	159	4	US-10-425-115-334563	Sequence 334563, A	370	30	65.2	501	4	US-10-425-115-26138	Sequence 26138, A
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299	30	65.2	168	4	US-10-424-599-197053	Sequence 197053, A	372	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
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308	30	65.2	217	4	US-10-156-761-14440	Sequence 14440, A	381	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
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311	30	65.2	286	4	US-10-369-493-821	Sequence 821, Ap	384	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
312	30	65.2	301	4	US-10-282-122A-68620	Sequence 68620, A	385	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
313	30	65.2	302	4	US-10-369-493-6656	Sequence 6656, Ap	386	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
314	30	65.2	302	5	US-10-774-355A-2386	Sequence 2386, Ap	387	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
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317	30	65.2	310	5	US-10-774-355A-1784	Sequence 1784, Ap	390	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
318	30	65.2	317	5	US-10-450-763-60569	Sequence 60569, Ap	391	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A
319	30	65.2	332	4	US-10-369-493-5269	Sequence 5269, Ap	392	30	65.2	502	6	US-10-424-599-220478	Sequence 220478, A

393	30	65.2	1250	3	US-09-769-736-10	Sequence 10, Appl	466	29	63.0	229	4	US-10-282-122A-71091	Sequence 71091, A
394	30	65.2	1281	5	US-10-732-923-12909	Sequence 12909, A	467	29	63.0	229	4	US-10-282-122A-71490	Sequence 71490, A
395	30	65.2	1293	4	US-10-697-036-41	Sequence 41, Appl	468	29	63.0	231	5	US-10-450-763-60093	Sequence 60093, A
396	30	65.2	1298	5	US-10-732-923-12907	Sequence 12907, A	469	29	63.0	232	4	US-10-437-963-118713	Sequence 118713, A
397	30	65.2	1298	5	US-10-732-923-12908	Sequence 12908, A	470	29	63.0	236	4	US-10-767-701-56113	Sequence 56113, A
398	30	65.2	1307	4	US-10-697-036-16	Sequence 16, Appl	471	29	63.0	237	4	US-10-724-972A-3781	Sequence 3781, Ap
399	30	65.2	1311	4	US-10-437-963-125427	Sequence 125427,	472	29	63.0	241	4	US-10-369-493-10503	Sequence 10503, A
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401	30	65.2	1315	4	US-10-697-036-13	Sequence 13, Appl	474	29	63.0	250	3	US-09-925-299-508	Sequence 508, Ap
402	30	65.2	1344	3	US-09-738-626-6868	Sequence 6868, Ap	475	29	63.0	250	4	US-10-102-806-508	Sequence 508, Ap
403	30	65.2	1434	5	US-10-494-672-220	Sequence 220, App	476	29	63.0	252	4	US-10-408-765A-2849	Sequence 2849, Ap
404	30	65.2	1460	4	US-10-425-115-214639	Sequence 214639,	477	29	63.0	254	4	US-10-424-599-183750	Sequence 183750,
405	30	65.2	1526	5	US-10-450-763-36475	Sequence 36475, A	478	29	63.0	275	4	US-10-425-114-65918	Sequence 65918, A
406	30	65.2	1526	5	US-10-450-763-37648	Sequence 37648, A	479	29	63.0	279	4	US-10-282-122A-57857	Sequence 57857, A
407	30	65.2	1554	5	US-10-450-763-37647	Sequence 37647, A	480	29	63.0	280	4	US-10-369-493-4531	Sequence 4531, Ap
408	30	65.2	1568	4	US-10-437-963-171037	Sequence 171037,	481	29	63.0	280	4	US-10-369-493-7289	Sequence 7289, Ap
409	30	65.2	1584	5	US-10-450-763-37649	Sequence 37649, A	482	29	63.0	284	5	US-10-450-763-37918	Sequence 37918, A
410	30	65.2	1627	5	US-10-450-763-36476	Sequence 36476, A	483	29	63.0	295	4	US-10-425-115-193656	Sequence 193656,
411	30	65.2	1807	4	US-10-437-963-128920	Sequence 128920,	484	29	63.0	297	4	US-10-288-160-8	Sequence 8,
412	30	65.2	1881	3	US-09-998-425-3	Sequence 3, Appl	485	29	63.0	301	4	US-10-424-599-235626	Sequence 235626,
413	30	65.2	1881	3	US-09-997-977-3	Sequence 3, Appl	486	29	63.0	301	4	US-10-424-599-235627	Sequence 235627,
414	30	65.2	2288	4	US-10-369-493-6774	Sequence 6774, Ap	487	29	63.0	301	4	US-10-424-599-235630	Sequence 235630,
415	30	65.2	3692	4	US-10-282-122A-71235	Sequence 71235, A	488	29	63.0	301	4	US-10-424-599-235633	Sequence 235633,
416	30	65.2	3696	4	US-10-724-972A-5942	Sequence 5942, Ap	489	29	63.0	306	4	US-10-437-963-159433	Sequence 159433,
417	30	65.2	4961	4	US-10-114-153-64	Sequence 64, Appl	490	29	63.0	309	4	US-10-072-012-569	Sequence 569, Ap
418	29	63.0	20	4	US-10-083-919A-27	Sequence 27, Appl	491	29	63.0	310	4	US-10-276-774-2674	Sequence 2674, Ap
419	29	63.0	20	4	US-10-083-919A-113	Sequence 113, App	492	29	63.0	312	5	US-10-739-930-6561	Sequence 6561, Ap
420	29	63.0	43	4	US-10-425-115-248110	Sequence 248110,	493	29	63.0	319	4	US-10-282-122A-76542	Sequence 76542, A
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422	29	63.0	50	3	US-09-864-761-48058	Sequence 48058, A	495	29	63.0	327	4	US-10-425-115-249040	Sequence 249040, A
423	29	63.0	54	4	US-10-425-115-301847	Sequence 301847, A	496	29	63.0	333	4	US-10-425-114-39744	Sequence 39744, A
424	29	63.0	57	4	US-10-424-599-153178	Sequence 153178,	497	29	63.0	334	4	US-10-369-493-21697	Sequence 21697, A
425	29	63.0	58	4	US-10-424-599-188899	Sequence 188899,	498	29	63.0	335	4	US-10-369-493-21864	Sequence 21864, A
426	29	63.0	67	4	US-10-424-599-250260	Sequence 250260,	499	29	63.0	335	4	US-10-451-467A-86	Sequence 486, Ap
427	29	63.0	73	4	US-10-437-963-201949	Sequence 201949,	500	29	63.0	337	5	US-10-282-122A-78237	Sequence 78237, A
428	29	63.0	76	4	US-10-424-599-257104	Sequence 257104,	501	29	63.0	338	4	US-10-450-763-31119	Sequence 31119, A
429	29	63.0	79	4	US-10-437-963-118212	Sequence 118212,	502	29	63.0	338	4	US-10-424-599-155317	Sequence 155317,
430	29	63.0	85	4	US-10-083-919A-190	Sequence 190, App	503	29	63.0	338	4	US-10-424-599-155322	Sequence 155322,
431	29	63.0	86	4	US-10-083-919A-26	Sequence 26, Appl	504	29	63.0	341	4	US-10-424-599-155322	Sequence 155318,
432	29	63.0	86	4	US-10-083-919A-90	Sequence 90, Appl	505	29	63.0	344	4	US-10-085-199-130	Sequence 130, Appl
433	29	63.0	86	4	US-10-083-919A-100	Sequence 100, App	506	29	63.0	345	4	US-10-200-012-20	Sequence 20, Appl
434	29	63.0	86	4	US-10-083-919A-183	Sequence 183, App	507	29	63.0	345	4	US-10-282-122A-49993	Sequence 49993, A
435	29	63.0	88	4	US-10-424-599-273308	Sequence 273308,	508	29	63.0	345	4	US-10-282-122A-49754	Sequence 49754, A
436	29	63.0	90	4	US-10-106-698-6200	Sequence 6200, Ap	509	29	63.0	347	4	US-10-423-114-45560	Sequence 45560, A
437	29	63.0	91	4	US-10-424-599-185601	Sequence 185601,	510	29	63.0	349	4	US-10-423-114-45560	Sequence 45560, A
438	29	63.0	98	4	US-10-767-701-38330	Sequence 38330, A	511	29	63.0	350	5	US-10-739-930-10621	Sequence 10621, A
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441	29	63.0	103	4	US-10-437-963-145205	Sequence 145205,	514	29	63.0	354	4	US-10-424-599-2133280	Sequence 213380,
442	29	63.0	119	4	US-10-108-260A-4479	Sequence 4479, Ap	515	29	63.0	355	4	US-10-425-114-56070	Sequence 56070, A
443	29	63.0	124	4	US-10-425-115-302166	Sequence 302166,	516	29	63.0	356	3	US-09-764-853-570	Sequence 570, App
444	29	63.0	124	4	US-10-424-599-230337	Sequence 230337,	517	29	63.0	357	4	US-10-425-114-37929	Sequence 37929, A
445	29	63.0	128	4	US-10-425-115-229957	Sequence 229957,	518	29	63.0	357	4	US-10-425-114-234745	Sequence 234745,
446	29	63.0	129	4	US-10-425-115-236015	Sequence 236015,	519	29	63.0	363	4	US-10-369-493-6647	Sequence 6647, A
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448	29	63.0	137	4	US-10-425-113-296306	Sequence 296306,	521	29	63.0	375	4	US-10-437-963-109989	Sequence 109989,
449	29	63.0	140	5	US-10-856-499-692	Sequence 692, App	522	29	63.0	377	4	US-10-437-963-110855	Sequence 110855,
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453	29	63.0	169	5	US-10-869-630-12	Sequence 12, Appl	526	29	63.0	415	4	US-10-282-122A-70442	Sequence 70442, A
454	29	63.0	169	5	US-10-883-760-68	Sequence 68, Appl	527	29	63.0	420	4	US-10-369-493-9503	Sequence 9503, Ap
455	29	63.0	172	4	US-10-425-115-263964	Sequence 263964,	528	29	63.0	420	5	US-10-450-763-43256	Sequence 43256, A
456	29	63.0	174	4	US-10-767-701-38243	Sequence 38243, A	529	29	63.0	420	5	US-10-450-763-45796	Sequence 45796, A
457	29	63.0	187	4	US-10-389-647-607	Sequence 607, Appl	530	29	63.0	422	5	US-10-450-763-51741	Sequence 51741, A
458	29	63.0	193	4	US-10-017-161-46	Sequence 46, Appl	531	29	63.0	424	4	US-10-369-493-9299	Sequence 9299, Ap
459	29	63.0	195	4	US-10-424-599-242050	Sequence 242050,	532	29	63.0	428	4	US-10-369-493-15489	Sequence 15489, A
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461	29	63.0	213	4	US-10-424-599-155315	Sequence 155315,	534	29	63.0	430	4	US-10-369-493-16240	Sequence 16240, A
462	29	63.0	214	4	US-10-424-599-235624	Sequence 235624,	535	29	63.0	431	5	US-10-739-930-11039	Sequence 11039, A
463	29	63.0	215	5	US-10-869-630-26	Sequence 26, Appl	536	29	63.0	432	5	US-10-739-930-11041	Sequence 11041, A
464	29	63.0	221	4	US-10-424-599-277200	Sequence 277200,	537	29	63.0	436	4	US-10-094-886-152	Sequence 152, App
465	29	63.0	224	4	US-10-424-599-257424	Sequence 257424,	538	29	63.0	436	5	US-10-732-923-89388	Sequence 8388, Ap

539	29	63.0	437	3	US-09-973-322-5	Sequence 11, Appl	612	29	63.0	2383	5	US-10-771-241-302	Sequence 302, App	
540	29	63.0	437	4	US-10-170-385-313	Sequence 313, App	613	29	63.0	2887	4	US-10-479-875-8	Sequence 8, Appl	
541	29	63.0	437	5	US-10-417-375-46	Sequence 46, Appl	614	29	63.0	3092	3	US-09-801-346-172	Sequence 172, Appl	
542	29	63.0	437	5	US-10-417-375-50	Sequence 50, Appl	615	29	63.0	3092	4	US-10-369-493-1470	Sequence 1470, Ap	
543	29	63.0	437	5	US-10-417-375-52	Sequence 52, Appl	616	29	63.0	3321	6	US-11-075-234-4	Sequence 4, Appl1	
544	29	63.0	437	5	US-10-417-375-54	Sequence 54, Appl	617	29	63.0	3494	5	US-10-473-127-697	Sequence 697, App	
545	29	63.0	438	4	US-10-338-777-55	Sequence 55, Appl	618	29	63.0	3494	5	US-10-473-127-703	Sequence 703, App	
546	29	63.0	449	4	US-10-282-122A-76636	Sequence 76636, A	619	29	63.0	3494	5	US-10-473-127-703	Sequence 703, App	
547	29	63.0	450	4	US-10-425-115-193654	Sequence 193654, A	620	29	63.0	3523	4	US-10-473-127-698	Sequence 698, App	
548	29	63.0	461	4	US-10-187-626-4551	Sequence 59, Appl	621	29	63.0	3523	5	US-10-473-127-701	Sequence 701, App	
549	29	63.0	462	4	US-10-437-963-129577	Sequence 129577, A	622	29	63.0	3623	5	US-10-473-127-702	Sequence 702, App	
550	29	63.0	472	4	US-10-424-599-165549	Sequence 165549, A	623	29	63.0	3623	5	US-10-473-127-702	Sequence 702, App	
551	29	63.0	476	6	US-11-097-143-26544	Sequence 26544, A	624	29	63.0	3623	5	US-10-473-127-702	Sequence 702, App	
552	29	63.0	478	4	US-10-425-114-58912	Sequence 58912, A	625	29	63.0	3623	5	US-10-473-127-702	Sequence 702, App	
553	29	63.0	480	4	US-10-425-114-61022	Sequence 61022, A	626	29	63.0	3623	5	US-10-473-127-702	Sequence 702, App	
554	29	63.0	481	6	US-11-097-143-8211	Sequence 8211, Ap	627	28.5	62.0	701	6	US-11-097-143-8211	Sequence 8211, Ap	
555	29	63.0	483	4	US-10-369-493-7991	Sequence 7991, Ap	628	28	60.9	28.5	29	3	US-09-864-761-35242	Sequence 35242, A
556	29	63.0	485	3	US-09-738-626-4551	Sequence 4551, Ap	629	28	60.9	38	3	US-09-864-761-35242	Sequence 35242, A	
557	29	63.0	502	4	US-10-296-115-1066	Sequence 1066, Ap	630	28	60.9	38	3	US-09-864-761-35242	Sequence 35242, A	
558	29	63.0	502	4	US-10-296-115-1066	Sequence 1066, Ap	631	28	60.9	45	4	US-10-425-115-187047	Sequence 187047, A	
559	29	63.0	502	4	US-10-437-963-198057	Sequence 198057, A	632	28	60.9	45	4	US-10-437-963-126975	Sequence 126975, A	
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561	29	63.0	511	4	US-10-437-963-192459	Sequence 192459, A	634	28	60.9	55	5	US-10-425-115-240317	Sequence 240317, A	
562	29	63.0	513	5	US-10-739-930-6164	Sequence 6164, Ap	635	28	60.9	58	4	US-10-424-599-230692	Sequence 230692, A	
563	29	63.0	530	4	US-10-211-060-3	Sequence 3, Appl1	636	28	60.9	59	4	US-10-424-599-230692	Sequence 230692, A	
564	29	63.0	530	5	US-10-631-467-1494	Sequence 1494, Ap	637	28	60.9	60	4	US-10-425-115-213240	Sequence 213240, A	
565	29	63.0	602	4	US-10-332-281-655	Sequence 655, App	638	28	60.9	61	4	US-10-425-115-203319	Sequence 203319, A	
566	29	63.0	612	6	US-11-097-143-4893	Sequence 4893, Ap	639	28	60.9	63	3	US-09-864-761-34870	Sequence 34870, A	
567	29	63.0	636	4	US-10-104-947-2449	Sequence 2449, Ap	640	28	60.9	63	3	US-09-864-761-34870	Sequence 34870, A	
568	29	63.0	636	5	US-10-450-763-35986	Sequence 35986, A	641	28	60.9	66	3	US-10-437-963-153893	Sequence 153893, A	
569	29	63.0	672	3	US-09-801-368-200	Sequence 200, App	642	28	60.9	66	3	US-10-437-963-153893	Sequence 153893, A	
570	29	63.0	693	4	US-10-297-895A-19	Sequence 20, Appl	643	28	60.9	67	4	US-10-424-599-178735	Sequence 178735, A	
571	29	63.0	701	4	US-10-297-895A-19	Sequence 19, Appl	644	28	60.9	68	4	US-10-424-599-178735	Sequence 178735, A	
572	29	63.0	705	4	US-10-108-360A-4346	Sequence 4346, Ap	645	28	60.9	68	4	US-10-424-599-178735	Sequence 178735, A	
573	29	63.0	726	4	US-10-424-599-262675	Sequence 262675, A	646	28	60.9	68	4	US-10-424-599-178735	Sequence 178735, A	
574	29	63.0	740	4	US-10-389-566-698	Sequence 698, App	647	28	60.9	69	5	US-10-424-599-178735	Sequence 178735, A	
575	29	63.0	761	4	US-10-225-486-55	Sequence 55, Appl	648	28	60.9	70	4	US-10-424-599-178735	Sequence 178735, A	
576	29	63.0	761	4	US-10-408-765A-696	Sequence 696, App	649	28	60.9	71	4	US-10-424-599-178735	Sequence 178735, A	
577	29	63.0	761	5	US-10-723-860-3785	Sequence 3785, Ap	650	28	60.9	74	4	US-10-425-115-264586	Sequence 264586, A	
578	29	63.0	761	5	US-10-756-149-5656	Sequence 5656, Ap	651	28	60.9	74	4	US-10-425-115-264586	Sequence 264586, A	
579	29	63.0	761	4	US-10-359-493-6493	Sequence 6493, Ap	652	28	60.9	76	4	US-10-424-599-172089	Sequence 172089, A	
580	29	63.0	766	4	US-10-287-895A-21	Sequence 21, Appl	653	28	60.9	76	4	US-10-424-599-172089	Sequence 172089, A	
581	29	63.0	787	4	US-10-128-714-8587	Sequence 8587, Ap	654	28	60.9	79	5	US-10-424-599-172089	Sequence 172089, A	
582	29	63.0	791	4	US-10-471-450-18	Sequence 18, Appl	655	28	60.9	80	3	US-09-864-761-48218	Sequence 48218, A	
583	29	63.0	792	4	US-10-282-122A-61193	Sequence 61193, A	656	28	60.9	80	3	US-10-424-599-192498	Sequence 192498, A	
584	29	63.0	815	5	US-10-450-763-37884	Sequence 37884, A	657	28	60.9	83	4	US-10-424-599-147254	Sequence 147254, A	
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586	29	63.0	821	4	US-10-171-404A-48	Sequence 48, Appl	659	28	60.9	92	4	US-10-312-829-15	Sequence 15, Appl	
587	29	63.0	829	4	US-10-437-963-156165	Sequence 156165, A	660	28	60.9	92	4	US-10-312-829-15	Sequence 1561, Ap	
588	29	63.0	850	3	US-09-904-389-2	Sequence 2, Appl1	661	28	60.9	92	4	US-10-312-829-15	Sequence 1561, Ap	
589	29	63.0	861	6	US-11-097-143-14901	Sequence 14901, A	662	28	60.9	93	4	US-10-312-829-15	Sequence 1561, Ap	
590	29	63.0	883	6	US-11-097-143-22419	Sequence 22419, A	663	28	60.9	93	4	US-10-312-829-15	Sequence 1561, Ap	
591	29	63.0	885	4	US-10-424-599-197333	Sequence 197333, A	664	28	60.9	94	5	US-10-424-599-166062	Sequence 166062, A	
592	29	63.0	891	4	US-10-408-765A-565	Sequence 565, Appl	665	28	60.9	94	5	US-10-424-599-166062	Sequence 166062, A	
593	29	63.0	927	4	US-10-297-895A-10	Sequence 10, Appl	666	28	60.9	95	4	US-10-424-599-203997	Sequence 203997, A	
594	29	63.0	949	4	US-10-437-963-121874	Sequence 121874, A	667	28	60.9	97	4	US-10-424-599-150518	Sequence 150518, A	
595	29	63.0	969	4	US-10-282-122A-76846	Sequence 76846, A	668	28	60.9	98	4	US-10-424-599-249338	Sequence 249338, A	
596	29	63.0	973	5	US-10-450-763-58453	Sequence 58453, A	669	28	60.9	99	4	US-10-424-599-249338	Sequence 249338, A	
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880	28	60.9	337	3	US-09-738-626-5808	Sequence 5808, App	953	28	60.9	411	5	US-10-450-763-34255	Sequence 34255, A
881	28	60.9	337	4	US-10-369-493-19085	Sequence 19085, A	954	28	60.9	412	5	US-10-450-763-43063	Sequence 43063, A
882	28	60.9	337	4	US-10-761-014-288	Sequence 288, App	955	28	60.9	415	4	US-10-081-816-6	Sequence 6, Appl1
883	28	60.9	337	5	US-10-494-836-60	Sequence 60, Appl1	956	28	60.9	417	4	US-10-437-963-139119	Sequence 139119, A
884	28	60.9	339	3	US-09-738-626-3344	Sequence 6344, Ap	957	28	60.9	417	5	US-10-437-963-34253	Sequence 34253, A
885	28	60.9	339	3	US-10-627-476-230	Sequence 230, App	958	28	60.9	418	5	US-10-450-763-33855	Sequence 33855, A
886	28	60.9	341	5	US-10-450-763-50239	Sequence 50239, A	959	28	60.9	419	5	US-10-450-763-33855	Sequence 38855, A
887	28	60.9	342	5	US-10-450-763-47690	Sequence 47690, A	960	28	60.9	421	4	US-10-425-115-355144	Sequence 355144, A
888	28	60.9	343	4	US-10-282-122A-54766	Sequence 54766, A	961	28	60.9	426	4	US-10-214-524-27	Sequence 27, Appl
889	28	60.9	343	5	US-10-450-763-11418	Sequence 41418, A	962	28	60.9	427	4	US-10-437-963-196346	Sequence 196346, A
890	28	60.9	347	5	US-10-450-763-11418	Sequence 47874, A	963	28	60.9	427	5	US-10-450-763-37855	Sequence 37855, A
891	28	60.9	348	5	US-10-450-763-9275	Sequence 39275, A	964	28	60.9	427	5	US-10-450-763-39001	Sequence 39001, A
892	28	60.9	348	5	US-10-450-763-57900	Sequence 57900, A	965	28	60.9	427	5	US-10-450-763-39001	Sequence 44856, A
893	28	60.9	349	5	US-10-450-763-39016	Sequence 39016, A	966	28	60.9	429	5	US-10-450-763-43435	Sequence 44856, A
894	28	60.9	350	4	US-10-437-963-161723	Sequence 161723, A	967	28	60.9	429	5	US-10-369-493-22639	Sequence 22639, A
895	28	60.9	351	4	US-10-282-122A-73666	Sequence 73666, A	968	28	60.9	430	4	US-10-450-763-33858	Sequence 64265, A
896	28	60.9	351	4	US-10-437-963-150161	Sequence 150161, A	969	28	60.9	431	5	US-10-450-763-33858	Sequence 33858, A
897	28	60.9	351	4	US-10-437-963-151726	Sequence 151726, A	970	28	60.9	432	5	US-10-450-763-51146	Sequence 51146, A
898	28	60.9	351	5	US-10-472-928-548	Sequence 548, App	971	28	60.9	433	5	US-10-450-763-51140	Sequence 51140, A
899	28	60.9	352	4	US-10-289-762-582	Sequence 582, App	972	28	60.9	438	4	US-10-062-848-2	Sequence 2, Appl1
900	28	60.9	352	4	US-10-437-963-120330	Sequence 120330, A	973	28	60.9	438	4	US-10-166-984-60	Sequence 60, Appl1
901	28	60.9	353	5	US-10-450-763-58440	Sequence 58440, A	974	28	60.9	438	4	US-10-166-984-60	Sequence 60, Appl1
902	28	60.9	353	5	US-10-450-763-36623	Sequence 36623, A	975	28	60.9	438	4	US-10-166-984-60	Sequence 60, Appl1
903	28	60.9	353	5	US-10-450-763-36957	Sequence 36957, A	976	28	60.9	438	4	US-10-776-104-2	Sequence 2, Appl1

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977 28 60.9 439 4 US-10-408-765A-2655 Sequence 2655, App
978 28 60.9 440 4 US-10-442-538-102 Sequence 102, App
979 28 60.9 440 4 US-10-442-538-125 Sequence 125, App
980 28 60.9 440 4 US-10-442-538-147 Sequence 147, App
981 28 60.9 440 4 US-10-492-782-20 Sequence 20, App
982 28 60.9 440 4 US-10-450-763-47887 Sequence 47887, A
983 28 60.9 441 4 US-10-177-293-146 Sequence 146, App
984 28 60.9 441 4 US-10-450-763-43408 Sequence 43408, A
985 28 60.9 441 4 US-10-450-763-56986 Sequence 56986, A
986 28 60.9 441 4 US-10-437-963-112507 Sequence 112507,
987 28 60.9 446 4 US-10-425-115-285083 Sequence 285083,
988 28 60.9 449 4 US-10-450-763-48348 Sequence 48348, A
989 28 60.9 450 4 US-10-062-848-15 Sequence 15, App
990 28 60.9 450 4 US-10-776-104-15 Sequence 15, App
991 28 60.9 452 4 US-10-450-763-34892 Sequence 34892, A
992 28 60.9 452 4 US-10-450-763-37868 Sequence 37868, A
993 28 60.9 452 4 US-10-450-763-44115 Sequence 44115, A
994 28 60.9 454 4 US-10-450-763-37286 Sequence 37286, A
995 28 60.9 454 4 US-10-450-763-38225 Sequence 38225, A
996 28 60.9 455 4 US-10-450-763-46091 Sequence 46091, A
997 28 60.9 456 4 US-10-474-792-144 Sequence 144, App
998 28 60.9 456 4 US-10-450-763-35471 Sequence 35471, A
999 28 60.9 456 4 US-10-450-763-43633 Sequence 43633, A
1000 28 60.9 457 4 US-10-437-963-117985 Sequence 117985,
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ALIGNMENTS

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RESULT 1
US-10-484-063-2
; Sequence 2, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILTERMO
; APPLICANT: FOLLEEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-2

Query Match          100.0%; Score 46; DB 5; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 2
US-10-476-570-21
; Sequence 21, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
```

```
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 21
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 20-34
US-10-476-570-21
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Query Match          100.0%; Score 46; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 3
US-10-476-570-22
; Sequence 22, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 22
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 24-38
US-10-476-570-22

Query Match          100.0%; Score 46; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.062;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 4
US-10-476-570-8
; Sequence 8, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
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; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIER, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 8
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 14-34
US-10-476-570-8

Query Match          100.0%; Score 46; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.089;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
DB 12 ELQTTTHDI 20

RESULT 5
US-11-021-949-1
; Sequence 1, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 1
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-1

Query Match          100.0%; Score 46; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
DB 17 ELQTTTHDI 25

RESULT 6
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
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; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIER, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 15-44
US-10-476-570-53

Query Match          100.0%; Score 46; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.13;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
DB 11 ELQTTTHDI 19

RESULT 7
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US20050033025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE B6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match          100.0%; Score 46; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.13;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
DB 11 ELQTTTHDI 19

RESULT 8
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US-10-476-570-9
; Sequence 9, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 32
; TYPE: PRF
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9
Query Match          100.0%; Score 46; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 12 ELQTTIHDI 20

RESULT 9
US-10-476-570-19
; Sequence 19, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 33
; TYPE: PRF
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19
Query Match          100.0%; Score 46; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
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Db 12 ELQTTIHDI 20

RESULT 10
US-10-177-390-6
; Sequence 6, Application US/10177390
; Publication No. US20030143743A1
; GENERAL INFORMATION:
; APPLICANT: Schuler, Gerold
; APPLICANT: N.V. Antwerp Innovatiecentrum
; TITLE OF INVENTION: Improved Transfection of Eucaryotic Cells with Linear
; FILE REFERENCE: 021505wo/JH/ml
; CURRENT APPLICATION NUMBER: US/10/177,390
; PRIOR FILING DATE: 2002-06-20
; NUMBER OF SEQ ID NOS: 34
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 151
; TYPE: PRF
; ORGANISM: Human papillomavirus type 16
US-10-177-390-6
Query Match          100.0%; Score 46; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.77;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 18 ELQTTIHDI 26

RESULT 11
US-10-484-063-20
; Sequence 20, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
; APPLICANT: TORTOLERO-LUNA, GUILLEMO
; APPLICANT: FOLLEN, MICHELLE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; PRIOR FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 151
; TYPE: PRF
; ORGANISM: Human papillomavirus
US-10-484-063-20
Query Match          100.0%; Score 46; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.77;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
Db 18 ELQTTIHDI 26

RESULT 12
US-10-484-063-27
; Sequence 27, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASTRY, K. JAGANNADHA
```



```

; APPLICANT: TORTOLERO-LUNA, GUILLERMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
; FILE REFERENCE: UTSC:560US
; CURRENT APPLICATION NUMBER: US/10/484,063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306,809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 27
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match          100.0%; Score 46; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.77;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 BLQTTTHDI 9
DB 18 BLQTTTHDI 26

RESULT 13
US-10-858-384-2
; Sequence 2, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match          100.0%; Score 46; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.81;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 BLQTTTHDI 9
DB 25 BLQTTTHDI 33

RESULT 14
US-10-367-057-16
; Sequence 16, Application US/10367057
; Publication No. US20050100554A1
; GENERAL INFORMATION:
; APPLICANT: Cuthill, Scott;
; APPLICANT: Jackson, Amanda;
; APPLICANT: Lewis, David A.;
; APPLICANT: Ooi, Chean Eng
; TITLE OF INVENTION: Complexes and Methods of Using Same
```

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; FILE REFERENCE: 21402-559
; CURRENT APPLICATION NUMBER: US/10/367,057
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: 60/256,911
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 198
; SOFTWARE: CuroSeqList version 0.1
; SEQ ID NO 16
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-367-057-16

Query Match          100.0%; Score 46; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.81;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 BLQTTTHDI 9
DB 25 BLQTTTHDI 33

RESULT 15
US-11-021-949-13
; Sequence 13, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SAMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match          100.0%; Score 46; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.81;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 BLQTTTHDI 9
DB 25 BLQTTTHDI 33

RESULT 16
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; CURRENT FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03271
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
```

SEQ ID NO 2
LENGTH: 171
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match 100.0%; Score 46; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 0.88;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
DB 30 ELQTTIHDI 38

RESULT 17
US-11-072-288-1
Sequence 1, Application US/11072288
Publication No. US20050159386A1
GENERAL INFORMATION:
APPLICANT: KIENY, Marie-Paule
APPLICANT: BALLOU, Jean-Marc
APPLICANT: BIZOUARNE, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
FILE REFERENCE: 01753-122
CURRENT APPLICATION NUMBER: US/11/072,288
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US/09/462,993
PRIOR FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: PCT/FR98/01576
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-11-072-288-1

Query Match 100.0%; Score 46; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 1.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
DB 53 ELQTTIHDI 61

RESULT 18
US-09-367-309A-1
Sequence 1, Application US/09367309A
Publication No. US20020081329A1
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALTIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6

SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 46; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
DB 25 ELQTTIHDI 33

RESULT 19
US-10-000-903-4
Sequence 4, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-4

Query Match 100.0%; Score 46; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
DB 131 ELQTTIHDI 139

RESULT 20
US-10-899-771-4
Sequence 4, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273

```
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: Influenzae B and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4
```

Query Match 100.0%; Score 46; DB 5; Length 273;

Best Local Similarity 100.0%; Pred. No. 1.5; Mismatches 0; Indels 0; Gaps 0;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy 1 ELQTTIHDI 9
    |||||
Db 131 ELQTTIHDI 139
```

RESULT 21

US-10-000-903-10

; Sequence 10, Application US/10000903

; Publication No. US20020182221A1

; GENERAL INFORMATION:

; APPLICANT: Bruck, Claudine

; APPLICANT: Cabezon Silva, Teresa

; APPLICANT: Delisse, Anne-Marie Eva Bernande

; APPLICANT: Gerard, Catherine Marie Ghislaine

; APPLICANT: Lombardo-Bencheikh, Angela

; TITLE OF INVENTION: Vaccine

; FILE REFERENCE: B45107

; CURRENT APPLICATION NUMBER: US/10/000, 903

; CURRENT FILING DATE: 2001-10-01

; PRIOR APPLICATION NUMBER: PCT/EP98/05285

; PRIOR FILING DATE: 1998-08-17

; PRIOR APPLICATION NUMBER: GB 9717953.5

; PRIOR FILING DATE: 1997-08-22

; NUMBER OF SEQ ID NOS: 23

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 10

; LENGTH: 292

; TYPE: PRT

; ORGANISM: Homo sapien

US-10-000-903-10

Query Match 100.0%; Score 46; DB 4; Length 292;

Best Local Similarity 100.0%; Pred. No. 1.6; Mismatches 0; Indels 0; Gaps 0;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy 1 ELQTTIHDI 9
    |||||
Db 150 ELQTTIHDI 158
```

RESULT 22

US-10-899-771-10

; Sequence 10, Application US/10899771

; Publication No. US20050031638A1

; GENERAL INFORMATION:

; APPLICANT: Dalemans, Wilfried L.J.

; APPLICANT: Gerard, Catherine Marie Ghislaine

; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins

; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide

; FILE REFERENCE: B45124

; CURRENT APPLICATION NUMBER: US/10/899, 771

; CURRENT FILING DATE: 2004-07-27

; PRIOR APPLICATION NUMBER: US/09/581, 976

; PRIOR FILING DATE: 2000-06-20

; PRIOR APPLICATION NUMBER: PCT/EP98/08563

; PRIOR FILING DATE: 1998-12-18

; PRIOR APPLICATION NUMBER: GB 9727262.9

; PRIOR FILING DATE: 1997-12-24

; NUMBER OF SEQ ID NOS: 28

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 10

LENGTH: 292

; TYPE: PRT

; ORGANISM: Artificial Sequence

; FEATURE:

; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus

; OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type

; OTHER INFORMATION: 16)

US-10-899-771-10

Query Match 100.0%; Score 46; DB 5; Length 292;

Best Local Similarity 100.0%; Pred. No. 1.6; Mismatches 0; Indels 0; Gaps 0;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy 1 ELQTTIHDI 9
    |||||
Db 150 ELQTTIHDI 158
```

RESULT 23

US-10-000-903-6

; Sequence 6, Application US/10000903

; Publication No. US20020182221A1

; GENERAL INFORMATION:

; APPLICANT: Bruck, Claudine

; APPLICANT: Cabezon Silva, Teresa

; APPLICANT: Delisse, Anne-Marie Eva Bernande

; APPLICANT: Gerard, Catherine Marie Ghislaine

; APPLICANT: Lombardo-Bencheikh, Angela

; TITLE OF INVENTION: Vaccine

; FILE REFERENCE: B45107

; CURRENT APPLICATION NUMBER: US/10/000, 903

; CURRENT FILING DATE: 2001-10-01

; PRIOR APPLICATION NUMBER: PCT/EP98/05285

; PRIOR FILING DATE: 1998-08-17

; PRIOR APPLICATION NUMBER: GB 9717953.5

; PRIOR FILING DATE: 1997-08-22

; NUMBER OF SEQ ID NOS: 23

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 6

; LENGTH: 371

; TYPE: PRT

; ORGANISM: Homo sapien

US-10-000-903-6

Query Match 100.0%; Score 46; DB 4; Length 371;

Best Local Similarity 100.0%; Pred. No. 2.1; Mismatches 0; Indels 0; Gaps 0;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```
Qy 1 ELQTTIHDI 9
    |||||
Db 131 ELQTTIHDI 139
```

RESULT 24

US-10-899-771-6

; Sequence 6, Application US/10899771

; Publication No. US20050031638A1

; GENERAL INFORMATION:

; APPLICANT: Dalemans, Wilfried L.J.

; APPLICANT: Gerard, Catherine Marie Ghislaine

; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins

; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide

; FILE REFERENCE: B45124

; CURRENT APPLICATION NUMBER: US/10/899, 771

; CURRENT FILING DATE: 2004-07-27

; PRIOR APPLICATION NUMBER: US/09/581, 976

; PRIOR FILING DATE: 2000-06-20

; PRIOR APPLICATION NUMBER: PCT/EP98/08563

; PRIOR FILING DATE: 1998-12-18

; PRIOR APPLICATION NUMBER: GB 9727262.9

; PRIOR FILING DATE: 1997-12-24

; NUMBER OF SEQ ID NOS: 28

; SOFTWARE: FastSeq for Windows Version 3.0

; SEQ ID NO 6

```
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenza B and B6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-6
```

```
Query Match          100.0%; Score 46; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 ELQTTTHDI 9
    |||||
Db 131 ELQTTTHDI 139
```

RESULT 25

```
US-10-000-903-14
; Sequence 14; Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-14
```

```
Query Match          100.0%; Score 46; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 2.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 ELQTTTHDI 9
    |||||
Db 150 ELQTTTHDI 158
```

RESULT 26

```
US-10-899-771-14
; Sequence 14; Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
```

```
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-14
```

```
Query Match          100.0%; Score 46; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 2.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 ELQTTTHDI 9
    |||||
Db 150 ELQTTTHDI 158
```

RESULT 27

```
US-10-367-095-10
; Sequence 10; Application US/10367095
; Publication No. US20030228696A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: NO. US20030228696A1et Insect Cell Line
; FILE REFERENCE: 44149-10SI
; CURRENT APPLICATION NUMBER: US/10/367,095
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-095-10
```

```
Query Match          100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 ELQTTTHDI 9
    |||||
Db 495 ELQTTTHDI 503
```

```
RESULT 28
US-10-368-046-10
; Sequence 10; Application US/10368046
```

```
/ Publication No. US20040063188A1
/ GENERAL INFORMATION:
/ APPLICANT: Robin A. Robinson
/ APPLICANT: Victoria Cioce
/ TITLE OF INVENTION: Method for Isolation and Purification of
/ TITLE OF INVENTION: Expressed Gene Products In Vitro
/ FILE REFERENCE: 44149-3US1
/ CURRENT APPLICATION NUMBER: US/10/368, 046
/ PRIOR FILING DATE: 2003-02-15
/ PRIOR APPLICATION NUMBER: US 60/356,119
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,161
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,118
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,133
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,157
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,135
/ PRIOR FILING DATE: 2002-02-14
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 13
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 10
/ LENGTH: 536
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-368-046-10

Query Match      100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy      1 ELQTTIHDI 9
Db      495 ELQTTIHDI 503

RESULT 29
US-10-367-367-10
/ Sequence 10, Application US/10367367
/ Publication No. US20040121465A1
/ GENERAL INFORMATION:
/ APPLICANT: Robin A. Robinson
/ TITLE OF INVENTION: Optimization of Gene Sequences of
/ TITLE OF INVENTION: Virus-Like Particles for Expression in Insect Cells
/ FILE REFERENCE: 44149-2US1
/ CURRENT APPLICATION NUMBER: US/10/367,367
/ PRIOR FILING DATE: 2003-02-15
/ PRIOR APPLICATION NUMBER: US 60/356,119
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,161
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,118
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,133
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,157
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,156
/ PRIOR FILING DATE: 2002-02-14
/ NUMBER OF SEQ ID NOS: 13
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 10
/ LENGTH: 536
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-918-337-10

Query Match      100.0%; Score 46; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 10
/ LENGTH: 536
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-367-367-10

Query Match      100.0%; Score 46; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy      1 ELQTTIHDI 9
Db      495 ELQTTIHDI 503

RESULT 30
US-10-918-337-10
/ Sequence 10, Application US/10918337
/ Publication No. US20050118191A1
/ GENERAL INFORMATION:
/ APPLICANT: NOVAVAX, INC., et al.
/ TITLE OF INVENTION: Optimization of Gene Sequences of
/ TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
/ FILE REFERENCE: 19065/2132
/ CURRENT APPLICATION NUMBER: US/10/918,337
/ PRIOR FILING DATE: 2004-08-13
/ PRIOR APPLICATION NUMBER: PCT/US03/04473
/ PRIOR FILING DATE: 2003-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,119
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,161
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,118
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,133
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,157
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,156
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,123
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,113
/ PRIOR FILING DATE: 2002-02-14
/ PRIOR APPLICATION NUMBER: US 60/356,154
/ PRIOR FILING DATE: 2002-02-14
/ Remaining Prior Application data removed - See File Wrapper or PALM.
/ NUMBER OF SEQ ID NOS: 13
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 10
/ LENGTH: 536
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-918-337-10

Query Match      100.0%; Score 46; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy      1 ELQTTIHDI 9
Db      495 ELQTTIHDI 503

RESULT 31
US-10-820-155-42
/ Sequence 42, Application US/10820155
/ Publication No. US20050137126A1
```

```
/ GENERAL INFORMATION:
/ APPLICANT: Natlgunny, A/S
/ APPLICANT: Natlgunny, Dietmar
/ APPLICANT: Jensenius, Jens Christian
/ APPLICANT: Kongerslev, Lelf
/ APPLICANT: Mathiesen, Finn
/ TITLE OF INVENTION: Treatment of SARS in individuals
/ FILE REFERENCE: P 774 US00
/ CURRENT APPLICATION NUMBER: US/10/820,155
/ CURRENT FILING DATE: 2004-04-08
/ NUMBER OF SEQ ID NOS: 137
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 42
/ LENGTH: 248
/ TYPE: PRT
/ ORGANISM: Canis familiaris
US-10-820-155-42

Query Match      91.3%; Score 42; DB 5; Length 248;
Best Local Similarity 77.8%; Pred. No. 7.6;
Matches 7; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      1 ELQTTIHD1 9
        |||||:|:|
Db      106 ELQTTIHD1 114

RESULT 32
US-10-820-155-57
/ Sequence 57, Application US/10820155
/ Publication No. US20050137126A1
/ GENERAL INFORMATION:
/ APPLICANT: Natlimmune A/S
/ APPLICANT: Weilgunny, Dietmar
/ APPLICANT: Jensenius, Jens Christian
/ APPLICANT: Kongerslev, Lelf
/ APPLICANT: Mathiesen, Finn
/ TITLE OF INVENTION: Treatment of SARS in individuals
/ FILE REFERENCE: P 774 US00
/ CURRENT APPLICATION NUMBER: US/10/820,155
/ CURRENT FILING DATE: 2004-04-08
/ NUMBER OF SEQ ID NOS: 137
/ SOFTWARE: PatentIn version 3.1
/ SEQ ID NO 57
/ LENGTH: 248
/ TYPE: PRT
/ ORGANISM: Canis familiaris
US-10-820-155-57

Query Match      91.3%; Score 42; DB 5; Length 248;
Best Local Similarity 77.8%; Pred. No. 7.6;
Matches 7; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      1 ELQTTIHD1 9
        |||||:|:|
Db      106 ELQTTIHD1 114

RESULT 33
US-09-909-460-102
/ Sequence 102, Application US/09909460
/ Publication No. US20020182258A1
/ GENERAL INFORMATION:
/ APPLICANT: Luneford, Lynn B.
/ APPLICANT: Putnam, David
/ APPLICANT: Hedley, Mary Lynn
/ TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY OF NUCLEIC
/ TITLE OF INVENTION: ACID
/ FILE REFERENCE: 08191/014001
/ CURRENT APPLICATION NUMBER: US/09/909,460
/ CURRENT FILING DATE: 2001-07-18
/ PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/321,346
/ PRIOR FILING DATE: EARLIER FILING DATE: 1999-05-27
```

```
/ NUMBER OF SEQ ID NOS: 114
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 102
/ LENGTH: 9
/ TYPE: PRT
/ ORGANISM: Hepatitis B virus
US-09-909-460-102

Query Match      89.1%; Score 41; DB 3; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      2 LQTTIHD1 9
        |||||
Db      1 LQTTIHD1 8

RESULT 34
US-09-872-836-102
/ Sequence 102, Application US/09872836
/ Publication No. US20040142475A1
/ GENERAL INFORMATION:
/ APPLICANT: Barman, Shilpa P.
/ APPLICANT: McKeever, Una
/ APPLICANT: Hedley, Mary Lynne
/ TITLE OF INVENTION: DELIVERY SYSTEMS FOR BIOACTIVE AGENTS
/ FILE REFERENCE: 08191-018001
/ CURRENT APPLICATION NUMBER: US/09/872,836
/ CURRENT FILING DATE: 2001-06-01
/ PRIOR APPLICATION NUMBER: US 60/208,830
/ PRIOR FILING DATE: 2000-06-02
/ NUMBER OF SEQ ID NOS: 120
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 102
/ LENGTH: 9
/ TYPE: PRT
/ ORGANISM: Homo sapiens
US-09-872-836-102

Query Match      89.1%; Score 41; DB 3; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      2 LQTTIHD1 9
        |||||
Db      1 LQTTIHD1 8

RESULT 35
US-10-133-210-279
/ Sequence 279, Application US/10133210
/ Publication No. US20030103964A1
/ GENERAL INFORMATION:
/ APPLICANT: Delisi, Charles
/ APPLICANT: Berzofsky, Jay
/ APPLICANT: Gulukota, Kamalakara
/ APPLICANT: Vaccaro, Dennis
/ APPLICANT: Wang, Zhiping
/ APPLICANT: Zhang, Chao
/ TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
/ TITLE OF INVENTION: COMPOSITIONS THEREOF
/ FILE REFERENCE: BU-035AX
/ CURRENT APPLICATION NUMBER: US/10/133,210
/ CURRENT FILING DATE: 2002-04-26
/ NUMBER OF SEQ ID NOS: 281
/ SOFTWARE: PatentIn Ver. 2.0
/ SEQ ID NO 279
/ LENGTH: 9
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-279
```

Query Match 89.1%; Score 41; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
DB 1 LQTTIHDI 8

RESULT 36

US-10-758-970-102
; Sequence 102, Application US/10758970
; Publication No. US20050037086A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Heu, Yung-Yueh
; TITLE OF INVENTION: CONTINUOUS-FLOW METHOD FOR PREPARING MICROPARTICLES
; FILE REFERENCE: 08191-012001
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: US/09/715,708A
; PRIOR FILING DATE: 2000-11-17
; PRIOR APPLICATION NUMBER: US 60/166,516
; PRIOR FILING DATE: 1999-11-19
; NUMBER OF SEQ ID NOS: 109
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 102
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human papilloma virus
US-10-758-970-102

Query Match 89.1%; Score 41; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
DB 1 LQTTIHDI 8

RESULT 37

US-10-751-845-56
; Sequence 56, Application US/10751845
; Publication No. US20050100928A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Urban, Robert G.
; APPLICANT: Chicz, Roman M.
; TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
; FILE REFERENCE: 08191-013001
; CURRENT APPLICATION NUMBER: US/10/751,845
; CURRENT FILING DATE: 2004-01-05
; PRIOR APPLICATION NUMBER: US/09/664,225
; PRIOR FILING DATE: 2000-08-18
; PRIOR APPLICATION NUMBER: US 60/169,846
; PRIOR FILING DATE: 1999-12-09
; PRIOR APPLICATION NUMBER: US 60/154,665
; PRIOR FILING DATE: 1999-09-16
; NUMBER OF SEQ ID NOS: 163
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 56
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papilloma virus
US-10-751-845-56

Query Match 89.1%; Score 41; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
DB 1 LQTTIHDI 8

RESULT 38

US-10-452-024-95
; Sequence 95, Application US/10452024
; Publication No. US20040013687A1
; GENERAL INFORMATION:
; APPLICANT: Simpson, Lance
; APPLICANT: Park, Jung-Beak
; APPLICANT: Maksymowich, Andrew
; TITLE OF INVENTION: Compositions and Methods For Trans epithelial Molecular Transport
; FILE REFERENCE: 9855-9611
; CURRENT APPLICATION NUMBER: US/10/452,024
; CURRENT FILING DATE: 2003-06-02
; PRIOR APPLICATION NUMBER: 60/384,949
; PRIOR FILING DATE: 2002-05-31
; NUMBER OF SEQ ID NOS: 188
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 95
; LENGTH: 1198
; TYPE: PRT
; ORGANISM: Clostridium botulinum
US-10-452-024-95

Query Match 80.4%; Score 37; DB 4; Length 1198;
Best Local Similarity 66.7%; Pred. No. 3.8e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
DB 323 KFGTIVHDI 331

RESULT 39

US-10-476-570-20
; Sequence 20, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: GUILLET, Jean-Gerard
; APPLICANT: POUVELLE-MORATILLE, Sandra
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: Papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 20
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 17-31
US-10-476-570-20

Query Match 78.3%; Score 36; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 5;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTIH 7
DB 9 ELQTTIH 15

```
RESULT 40
US-10-425-115-237192
; Sequence 237192, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovacic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 237192
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(151)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_147905C.1.pep
US-10-425-115-237192
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```
Query Match          78.3%; Score 36; DB 4; Length 151;
Best Local Similarity 75.0%; Pred. No. 62;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 ELQTHDI 9
Db      96 ELQTHDI 103
```

```
RESULT 41
US-11-097-143-21504
; Sequence 21504, Application US/11097143
; Publication No. US20050208558A1
; GENERAL INFORMATION:
; APPLICANT: Venter, J. Craig
; APPLICANT: et al.
; TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
; TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
; TITLE OF INVENTION: DROSOPHILA GENES.
; FILE REFERENCE: CL000728
; CURRENT APPLICATION NUMBER: US/11/097,143
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: 60/157,832
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: 60/160,191
; PRIOR FILING DATE: 1999-10-19
; PRIOR APPLICATION NUMBER: 60/161,932
; PRIOR FILING DATE: 1999-10-28
; PRIOR APPLICATION NUMBER: 60/164,769
; PRIOR FILING DATE: 1999-11-12
; PRIOR APPLICATION NUMBER: 60/173,383
; PRIOR FILING DATE: 1999-12-28
; PRIOR APPLICATION NUMBER: 60/175,693
; PRIOR FILING DATE: 2000-01-12
; PRIOR APPLICATION NUMBER: 60/184,831
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60/191,637
; PRIOR FILING DATE: 2000-03-23
; NUMBER OF SEQ ID NOS: 43008
; SOFTWARE: FaastSeq for Windows Version 4.0
; SEQ ID NO 21504
; LENGTH: 194
; TYPE: PRT
; ORGANISM: DROSOPHILA
```

US-11-097-143-21504

```
Query Match          78.3%; Score 36; DB 6; Length 194;
Best Local Similarity 66.7%; Pred. No. 81;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTHDI 9
Db      55 ELQTHDI 63
```

```
RESULT 42
US-10-312-829-6
; Sequence 6, Application US/10312829
; Publication No. US20040037781A1
; GENERAL INFORMATION:
; APPLICANT: McCormack, Francis X
; TITLE OF INVENTION: Peptides with Antioxidant and Antimicrobial Properties
; FILE REFERENCE: 10738-31
; CURRENT APPLICATION NUMBER: US/10/312,829
; CURRENT FILING DATE: 2003-08-04
; PRIOR APPLICATION NUMBER: PCT/US01/21226
; PRIOR FILING DATE: 2001-06-29
; PRIOR APPLICATION NUMBER: US 60/215,313
; PRIOR FILING DATE: 2000-06-30
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 6
; LENGTH: 93
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-312-829-6
```

```
Query Match          76.1%; Score 35; DB 4; Length 93;
Best Local Similarity 75.0%; Pred. No. 56;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTHDI 8
Db      6 ELQTHDI 13
```

```
RESULT 43
US-09-925-302-845
; Sequence 845, Application US/09925302
; Patent No. US2002004941A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA104
; CURRENT APPLICATION NUMBER: US/09/925,302
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05918
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 896
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 845
; LENGTH: 145
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-925-302-845
```

```
Query Match          76.1%; Score 35; DB 3; Length 145;
Best Local Similarity 75.0%; Pred. No. 91;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 ELQTHDI 8
Db      123 ELQTHDI 130
```



```
RESULT 44
US-09-925-302-845
; Sequence 845, Application US/09925302
; Publication No. US20030064072A9
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins and Antibodies
; FILE REFERENCE: PA104
; CURRENT APPLICATION NUMBER: US/09/925,302
; CURRENT FILING DATE: 2001-08-10
; PRIOR APPLICATION NUMBER: PCT/US00/05918
; PRIOR FILING DATE: 2000-03-08
; PRIOR APPLICATION NUMBER: 60/124,270
; PRIOR FILING DATE: 1999-03-12
; NUMBER OF SEQ ID NOS: 896
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 845
; LENGTH: 145
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-925-302-845

Query Match      76.1%; Score 35; DB 3; Length 145;
Best Local Similarity 75.0%; Pred. No. 91;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      1 ELQTTIHD 8
Db      123 ELQATLHD 130

RESULT 45
US-10-312-829-5
; Sequence 5, Application US/10312829
; Publication No. US20040037781A1
; GENERAL INFORMATION:
; APPLICANT: McCormack, Francis X
; TITLE OF INVENTION: Peptides with Antioxidant and Antimicrobial Properties
; FILE REFERENCE: 10738-31
; CURRENT APPLICATION NUMBER: US/10/312,829
; CURRENT FILING DATE: 2003-08-04
; PRIOR APPLICATION NUMBER: PCT/US01/21226
; PRIOR FILING DATE: 2001-06-29
; PRIOR APPLICATION NUMBER: US 60/215,313
; PRIOR FILING DATE: 2000-06-30
; NUMBER OF SEQ ID NOS: 17
; SOFTWARE: Patentin version 3.1
; SEQ ID NO 5
; LENGTH: 148
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-312-829-5

Query Match      76.1%; Score 35; DB 4; Length 148;
Best Local Similarity 75.0%; Pred. No. 93;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      1 ELQTTIHD 8
Db      6 ELQATLHD 13

RESULT 46
US-10-336-603A-98
; Sequence 98, Application US/10336603A
; Publication No. US20040072997A1
; GENERAL INFORMATION:
; APPLICANT: Alsobrook et al.
; TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD
; FILE REFERENCE: 21402-533A
; CURRENT APPLICATION NUMBER: US/10/336,603A
; CURRENT FILING DATE: 2003-01-03
; PRIOR APPLICATION NUMBER: 09/746,491
```

```
; PRIOR FILING DATE: 2000-12-20
; PRIOR APPLICATION NUMBER: 10/055,569
; PRIOR FILING DATE: 2001-10-26
; NUMBER OF SEQ ID NOS: 169
; SOFTWARE: Curaseqdist version 0.1
; SEQ ID NO 98
; LENGTH: 166
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-336-603A-98

Query Match      76.1%; Score 35; DB 4; Length 166;
Best Local Similarity 75.0%; Pred. No. 1,1e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      1 ELQTTIHD 8
Db      24 ELQATLHD 31

RESULT 47
US-10-437-963-163049
; Sequence 163049, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovacic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barzak, Brad
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 163049
; LENGTH: 233
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_62080C.1.pep
US-10-437-963-163049

Query Match      76.1%; Score 35; DB 4; Length 233;
Best Local Similarity 62.5%; Pred. No. 1.5e+02;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy      1 ELQTTIHD 8
Db      186 EMQSTVHD 193

RESULT 48
US-10-336-603A-96
; Sequence 96, Application US/10336603A
; Publication No. US20040072997A1
; GENERAL INFORMATION:
; APPLICANT: Alsobrook et al.
; TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD
; FILE REFERENCE: 21402-533A
; CURRENT APPLICATION NUMBER: US/10/336,603A
; CURRENT FILING DATE: 2003-01-03
; PRIOR APPLICATION NUMBER: 09/746,491
; PRIOR FILING DATE: 2000-12-20
; PRIOR APPLICATION NUMBER: 10/055,569
; PRIOR FILING DATE: 2001-10-26
; NUMBER OF SEQ ID NOS: 169
; SOFTWARE: Curaseqdist version 0.1
; SEQ ID NO 96
```

LENGTH: 243
TYPE: PRT
ORGANISM: Homo sapiens
US-10-336-603A-96

Query Match 76.1%; Score 35; DB 4; Length 243;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
||| : ||
Db 101 ELQATLHD 108

RESULT 49
US-10-312-829-4
Sequence 4, Application US/10312829
Publication No. US2004003781A1
GENERAL INFORMATION:
APPLICANT: McCormack, Francis X
TITLE OF INVENTION: Peptides with Antioxidant and Antimicrobial Properties
FILE REFERENCE: 10738-31
CURRENT APPLICATION NUMBER: US/10/312,829
CURRENT FILING DATE: 2003-08-04
PRIOR APPLICATION NUMBER: PCT/US01/21226
PRIOR FILING DATE: 2001-06-29
PRIOR APPLICATION NUMBER: US 60/215,313
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 17
SOFTWARE: PatentIn version 3.1
SEQ ID NO 4
LENGTH: 248
TYPE: PRT
ORGANISM: Homo sapiens
US-10-312-829-4

Query Match 76.1%; Score 35; DB 4; Length 248;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
||| : ||
Db 106 ELQATLHD 113

RESULT 50
US-10-336-603A-100
Sequence 100, Application US/10336603A
Publication No. US20040072997A1
GENERAL INFORMATION:
APPLICANT: Alsbrook et al.
TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD
FILE REFERENCE: 21402-533A
CURRENT APPLICATION NUMBER: US/10/336,603A
CURRENT FILING DATE: 2003-01-03
PRIOR APPLICATION NUMBER: 09/746,491
PRIOR FILING DATE: 2000-12-20
PRIOR APPLICATION NUMBER: 10/055,569
PRIOR FILING DATE: 2001-10-26
NUMBER OF SEQ ID NOS: 169
SOFTWARE: Curaseqdist version 0.1
SEQ ID NO 100
LENGTH: 248
TYPE: PRT
ORGANISM: Homo sapiens
US-10-336-603A-100

Query Match 76.1%; Score 35; DB 4; Length 248;
Best Local Similarity 75.0%; Pred. No. 1.6e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
||| : ||

Db 106 ELQATLHD 113
Search completed: May 5, 2006, 08:07:21
Job time : 61 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 07:56:56 ; Search time 8.4 Seconds
(without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-5
Perfect score: 46
Sequence: 1 ELQRTTHDI 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

1: /SIDS5/ptodata/1/pubpaa/US08_NEW_PUB.pep1:*
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3: /SIDS5/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
4: /SIDS5/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
5: /SIDS5/ptodata/1/pubpaa/PCR_NEW_PUB.pep:*
6: /SIDS5/ptodata/1/pubpaa/US03_NEW_PUB.pep:*
7: /SIDS5/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
8: /SIDS5/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
9: /SIDS5/ptodata/1/pubpaa/US11_NEW_PUB.pep1:*
10: /SIDS5/ptodata/1/pubpaa/US11_NEW_PUB.pep1:*
11: /SIDS5/ptodata/1/pubpaa/US11_NEW_PUB.pep1:*
12: /SIDS5/ptodata/1/pubpaa/US60_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	46	100.0	151	9	US-10-530-253-13
2	46	100.0	158	11	US-11-206-138-3
3	46	100.0	248	9	US-10-530-253-1
4	46	100.0	248	9	US-10-530-253-3
5	46	100.0	248	9	US-10-530-253-5
6	46	100.0	248	9	US-10-530-253-7
7	46	100.0	248	9	US-10-530-253-9
8	46	100.0	248	9	US-10-530-253-11
9	46	100.0	256	11	US-11-092-923A-2
10	46	87.0	250	11	US-11-096-568A-1943
11	40	87.0	298	11	US-11-096-568A-1942
12	40	87.0	303	11	US-11-096-568A-1941
13	35	76.1	19	9	US-10-503-575-174
14	35	76.1	133	9	US-10-986-405-199
15	35	76.1	135	9	US-10-986-405-205
16	35	76.1	140	9	US-10-986-405-229
17	35	76.1	149	9	US-10-986-405-200
18	35	76.1	184	9	US-10-986-405-216
19	34	73.9	287	11	US-11-096-568A-19577
20	34	73.9	383	9	US-10-703-799B-62
21	34	73.9	448	11	US-11-096-568A-19576

22	34	73.9	463	11	US-11-096-568A-19575	Sequence 19575, A
23	34	73.9	628	11	US-11-080-991-108	Sequence 108, App
24	34	73.9	1479	9	US-10-204-639-4	Sequence 4, Appl1
25	34	73.9	1896	9	US-10-877-346-13	Sequence 13, Appl1
26	34	73.9	1905	9	US-10-877-346-44	Sequence 44, Appl1
27	33	71.7	149	9	US-10-530-253-17	Sequence 17, Appl1
28	33	71.7	158	9	US-10-530-253-26	Sequence 26, Appl1
29	33	71.7	237	9	US-10-506-454-1603	Sequence 1603, App
30	33	71.7	1763	8	US-10-504-120-21	Sequence 21, Appl1
31	33	71.7	1807	8	US-10-504-120-22	Sequence 22, Appl1
32	32	69.6	162	11	US-11-087-059-2617	Sequence 2617, App
33	32	69.6	333	11	US-11-188-298-17532	Sequence 17532, A
34	32	69.6	337	11	US-11-188-298-16137	Sequence 16137, A
35	32	69.6	340	11	US-11-188-298-6731	Sequence 6731, App
36	32	69.6	498	11	US-11-188-298-3477	Sequence 3477, App
37	32	69.6	513	9	US-10-873-528-37	Sequence 37, Appl1
38	32	69.6	513	9	US-10-873-528-193	Sequence 193, App
39	32	69.6	553	11	US-11-103-957-61	Sequence 193, App
40	32	69.6	643	11	US-11-087-059-1991	Sequence 1991, App
41	32	69.6	989	11	US-11-188-298-5563	Sequence 5563, App
42	31	67.4	329	11	US-11-096-568A-6040	Sequence 6040, App
43	31	67.4	335	11	US-11-188-298-21765	Sequence 21765, A
44	31	67.4	340	11	US-11-188-298-16409	Sequence 16409, A
45	31	67.4	540	11	US-11-096-568A-6039	Sequence 6039, App
46	31	67.4	917	11	US-11-169-041-145	Sequence 145, App
47	31	67.4	1039	9	US-10-915-002-321	Sequence 321, App
48	31	67.4	1039	9	US-10-915-002-322	Sequence 321, App
49	31	67.4	1039	9	US-10-915-002-323	Sequence 323, App
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51	31	67.4	1473	11	US-11-096-568A-14691	Sequence 14691, A
52	31	67.4	1529	11	US-11-096-568A-14690	Sequence 14690, A
53	30	65.2	63	11	US-11-079-463-6103	Sequence 6103, App
54	30	65.2	228	9	US-10-467-657-2280	Sequence 2280, App
55	30	65.2	293	11	US-11-045-004-1209	Sequence 1209, App
56	30	65.2	291	11	US-11-188-298-17244	Sequence 17244, A
57	30	65.2	333	11	US-11-188-298-3948	Sequence 3948, App
58	30	65.2	333	11	US-11-188-298-22053	Sequence 22053, A
59	30	65.2	334	11	US-11-188-298-2051	Sequence 2051, App
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61	30	65.2	348	11	US-11-087-059-2207	Sequence 2207, App
62	30	65.2	348	11	US-11-045-004-602	Sequence 602, App
63	30	65.2	352	11	US-11-188-298-12429	Sequence 12429, A
64	30	65.2	371	11	US-11-079-463-6883	Sequence 6883, App
65	30	65.2	373	11	US-11-045-004-6575	Sequence 6575, App
66	30	65.2	419	11	US-11-096-568A-21864	Sequence 21864, A
67	30	65.2	470	11	US-11-079-463-6501	Sequence 6501, App
68	30	65.2	474	11	US-11-096-568A-21863	Sequence 21863, A
69	30	65.2	876	11	US-11-045-004-266	Sequence 266, App
70	30	65.2	1250	11	US-11-123-241-10	Sequence 1142, App
71	30	65.2	4913	9	US-10-453-372-1142	Sequence 1142, App
72	30	65.2	4961	9	US-10-453-372-1132	Sequence 1132, App
73	29	63.0	15	9	US-10-530-061-1673	Sequence 1673, App
74	29	63.0	149	9	US-10-530-253-24	Sequence 2528, App
75	29	63.0	155	9	US-10-793-626-2528	Sequence 2528, App
76	29	63.0	159	11	US-11-183-664-112	Sequence 12, Appl1
77	29	63.0	184	11	US-11-188-298-2828	Sequence 2828, App
78	29	63.0	195	11	US-11-188-298-578	Sequence 578, App
79	29	63.0	196	11	US-11-087-059-8359	Sequence 8359, App
80	29	63.0	197	11	US-11-087-059-4852	Sequence 4852, App
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83	29	63.0	215	11	US-11-183-664-26	Sequence 26, Appl1
84	29	63.0	242	11	US-11-188-298-22212	Sequence 22212, App
85	29	63.0	243	11	US-11-045-004-1371	Sequence 1371, App
86	29	63.0	295	11	US-11-188-298-12304	Sequence 12304, A
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93	29	63.0	331	11	US-11-188-298-15348	Sequence 15348, A
94	29	63.0	331	11	US-11-188-298-15995	Sequence 15995, A

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117	29	63.0	340	11	US-11-188-298-2839	Sequence 2839, Ap
118	29	63.0	340	11	US-11-188-298-6179	Sequence 6179, Ap
119	29	63.0	340	11	US-11-188-298-8037	Sequence 8037, Ap
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129	29	63.0	434	11	US-11-087-099-3939	Sequence 3939, Ap
130	29	63.0	437	9	US-10-821-234-1392	Sequence 1392, Ap
131	29	63.0	475	11	US-11-087-099-1613	Sequence 1613, Ap
132	29	63.0	636	11	US-11-072-512-2449	Sequence 2449, Ap
133	29	63.0	693	11	US-11-189-301-20	Sequence 20, Appl
134	29	63.0	701	11	US-11-189-301-19	Sequence 19, Appl
135	29	63.0	761	9	US-10-501-035-380	Sequence 380, App
136	29	63.0	766	11	US-11-189-301-21	Sequence 21, Appl
137	29	63.0	927	11	US-11-189-301-10	Sequence 10, Appl
138	29	63.0	3623	9	US-10-995-561-593	Sequence 593, App
139	28	60.9	102	9	US-10-816-768-46	Sequence 46, Appl
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141	28	60.9	134	11	US-11-096-568A-20906	Sequence 20906, A
142	28	60.9	139	11	US-11-188-298-15597	Sequence 15597, A
143	28	60.9	153	11	US-11-096-568A-20905	Sequence 20905, A
144	28	60.9	159	11	US-11-087-099-1865	Sequence 1865, Ap
145	28	60.9	163	9	US-10-506-454-823	Sequence 823, App
146	28	60.9	164	11	US-11-087-099-7954	Sequence 7954, Ap
147	28	60.9	170	11	US-11-079-463-9969	Sequence 9969, Ap
148	28	60.9	181	11	US-11-264-096-910	Sequence 910, App
149	28	60.9	186	11	US-11-087-099-10391	Sequence 10391, A
150	28	60.9	187	11	US-11-087-099-7508	Sequence 7508, Ap
151	28	60.9	202	11	US-11-087-099-1322	Sequence 1322, Ap
152	28	60.9	207	11	US-11-045-004-375	Sequence 375, App
153	28	60.9	209	11	US-11-096-568A-20904	Sequence 20904, A
154	28	60.9	260	9	US-10-506-454-1219	Sequence 1219, Ap
155	28	60.9	294	11	US-11-098-686-10545	Sequence 10545, A
156	28	60.9	304	11	US-11-098-686-10425	Sequence 10425, A
157	28	60.9	306	11	US-11-156-084-305	Sequence 305, App
158	28	60.9	311	11	US-11-096-568A-23505	Sequence 23505, A
159	28	60.9	311	11	US-11-106-715-79	Sequence 79, Appl
160	28	60.9	317	11	US-11-082-389-28	Sequence 28, Appl
161	28	60.9	319	11	US-11-184-005-8	Sequence 8, Appl
162	28	60.9	325	11	US-11-184-005-2	Sequence 4, Appl
163	28	60.9	325	11	US-11-184-005-4	Sequence 4, Appl
164	28	60.9	328	11	US-11-098-686-10153	Sequence 10153, A
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171	28	60.9	340	11	US-11-188-298-14766	Sequence 14766, A
172	28	60.9	341	11	US-11-188-298-18458	Sequence 18458, A
173	28	60.9	343	11	US-11-079-463-8668	Sequence 8668, Ap
174	28	60.9	345	9	US-10-467-657-252	Sequence 252, App
175	28	60.9	345	9	US-10-467-657-3086	Sequence 3086, Ap
176	28	60.9	350	11	US-11-096-568A-31866	Sequence 31866, A
177	28	60.9	351	11	US-11-098-686-10666	Sequence 10666, A
178	28	60.9	354	9	US-10-878-558A-84	Sequence 84, Appl
179	28	60.9	354	11	US-11-100-640-4	Sequence 4, Appl
180	28	60.9	360	9	US-10-650-328B-8	Sequence 8, Appl
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182	28	60.9	375	11	US-11-172-740-665	Sequence 665, App
183	28	60.9	382	11	US-11-096-568A-30018	Sequence 30018, A
184	28	60.9	400	11	US-11-096-568A-30017	Sequence 30017, A
185	28	60.9	464	11	US-11-087-099-4021	Sequence 4021, Ap
186	28	60.9	474	11	US-11-188-298-14119	Sequence 14119, A
187	28	60.9	475	11	US-11-087-099-1092	Sequence 1092, Ap
188	28	60.9	475	11	US-11-087-099-10530	Sequence 10530, A
189	28	60.9	486	11	US-11-072-512-3837	Sequence 3837, Ap
190	28	60.9	500	11	US-11-098-686-10274	Sequence 10274, Ap
191	28	60.9	540	11	US-11-188-298-316	Sequence 316, App
192	28	60.9	550	11	US-11-072-512-3416	Sequence 3416, Ap
193	28	60.9	557	9	US-10-512-109-9	Sequence 9, Appl
194	28	60.9	594	11	US-11-052-554A-379	Sequence 379, App
195	28	60.9	606	11	US-11-079-463-9950	Sequence 9950, Ap
196	28	60.9	666	11	US-11-188-298-18205	Sequence 18205, A
197	28	60.9	704	11	US-11-079-463-8082	Sequence 8082, A
198	28	60.9	725	11	US-11-078-189-15	Sequence 15, Appl
199	28	60.9	745	9	US-10-995-561-659	Sequence 659, App
200	28	60.9	745	11	US-11-109-156-14	Sequence 14, Appl
201	28	60.9	745	11	US-11-222-158-4	Sequence 4, Appl
202	28	60.9	778	11	US-11-264-096-912	Sequence 912, App
203	28	60.9	820	9	US-10-330-773-533	Sequence 533, App
204	28	60.9	847	9	US-10-455-772-282	Sequence 282, App
205	28	60.9	908	11	US-11-087-099-11020	Sequence 11020, A
206	28	60.9	967	8	US-10-505-928-795	Sequence 795, App
207	28	60.9	967	11	US-11-054-281-74	Sequence 74, Appl
208	28	60.9	1075	11	US-11-100-640-12	Sequence 12, Appl
209	28	60.9	1148	11	US-11-110-082-29	Sequence 29, Appl
210	28	60.9	1245	11	US-11-110-082-30	Sequence 30, Appl
211	28	60.9	1279	9	US-10-821-234-1598	Sequence 1598, Ap
212	28	60.9	1299	11	US-11-169-041-231	Sequence 231, App
213	28	60.9	1345	11	US-11-052-554A-282	Sequence 282, App
214	28	60.9	3028	9	US-10-455-772-276	Sequence 276, App
215	28	60.9	3028	9	US-10-455-772-284	Sequence 284, App
216	28	60.9	3028	9	US-10-455-772-286	Sequence 286, App
217	28	60.9	3028	9	US-10-455-772-288	Sequence 288, App
218	27	58.7	9	11	US-11-171-365-2	Sequence 2, Appl
219	27	58.7	26	9	US-10-510-246-46	Sequence 46, Appl
220	27	58.7	90	11	US-11-079-463-7737	Sequence 6737, Ap
221	27	58.7	95	11	US-11-188-298-19355	Sequence 19355, A
222	27	58.7	142	11	US-11-079-463-8099	Sequence 8099, Ap
223	27	58.7	153	11	US-11-188-298-6658	Sequence 6658, Ap
224	27	58.7	158	9	US-10-530-253-15	Sequence 15, Appl
225	27	58.7	161	11	US-11-045-004-455	Sequence 455, App
226	27	58.7	179	11	US-11-188-298-10295	Sequence 10295, A
227	27	58.7	190	11	US-11-188-298-18763	Sequence 18763, A
228	27	58.7	192	11	US-11-188-298-7671	Sequence 7671, Ap
229	27	58.7	196	11	US-11-188-298-5361	Sequence 5361, Ap
230	27	58.7	196	11	US-11-188-298-17112	Sequence 17112, A
231	27	58.7	206	11	US-11-087-099-2337	Sequence 2337, Ap
232	27	58.7	209	11	US-11-188-298-14162	Sequence 14162, A
233	27	58.7	212	11	US-11-188-298-7543	Sequence 7543, Ap
234	27	58.7	219	11	US-11-188-298-17133	Sequence 17133, A
235	27	58.7	234	11	US-11-188-298-20885	Sequence 20885, A
236	27	58.7	242	9	US-10-467-657-806	Sequence 806, App
237	27	58.7	245	11	US-11-188-298-15164	Sequence 15164, A
238	27	58.7	245	11	US-11-188-298-22452	Sequence 22452, A
239	27	58.7	247	11	US-11-188-298-20584	Sequence 20584, A
240	27	58.7	248	11	US-11-188-298-21697	Sequence 21697, A

241	27	58.7	256	11	US-11-087-099-5474	Sequence 5474, Ap	314	27	58.7	338	11	US-11-188-298-8233	Sequence 8233, Ap
242	27	58.7	252	11	US-11-087-099-9660	Sequence 9660, Ap	315	27	58.7	338	11	US-11-188-298-10456	Sequence 10456, A
243	27	58.7	253	11	US-11-087-099-5632	Sequence 5632, A	316	27	58.7	338	11	US-11-188-298-13405	Sequence 13405, A
244	27	58.7	263	11	US-11-087-099-10938	Sequence 10938, A	317	27	58.7	338	11	US-11-188-298-19825	Sequence 19825, A
245	27	58.7	267	11	US-11-188-298-22153	Sequence 22153, A	318	27	58.7	338	11	US-11-188-298-21341	Sequence 21341, A
246	27	58.7	278	11	US-11-188-298-5866	Sequence 5866, Ap	319	27	58.7	338	11	US-11-188-298-22226	Sequence 22226, A
247	27	58.7	279	9	US-10-506-454-1033	Sequence 1033, Ap	320	27	58.7	339	11	US-11-188-298-7132	Sequence 7132, Ap
248	27	58.7	280	11	US-11-096-568A-12916	Sequence 12916, A	321	27	58.7	339	11	US-11-188-298-15158	Sequence 15158, A
249	27	58.7	280	11	US-11-172-740-2144	Sequence 2144, A	322	27	58.7	339	11	US-11-188-298-4766	Sequence 4766, Ap
250	27	58.7	284	11	US-11-188-298-19079	Sequence 19079, A	323	27	58.7	340	11	US-11-188-298-8513	Sequence 8513, Ap
251	27	58.7	287	11	US-11-096-568A-12915	Sequence 12915, A	324	27	58.7	340	11	US-11-188-298-15728	Sequence 15728, A
252	27	58.7	303	11	US-11-188-298-4964	Sequence 4964, Ap	325	27	58.7	340	11	US-11-188-298-18968	Sequence 18968, A
253	27	58.7	305	11	US-11-188-298-13505	Sequence 13505, A	326	27	58.7	340	11	US-11-188-298-18968	Sequence 18968, A
254	27	58.7	310	11	US-11-188-298-7885	Sequence 7885, Ap	327	27	58.7	341	9	US-10-793-626-1188	Sequence 1188, Ap
255	27	58.7	315	11	US-11-188-298-11717	Sequence 11717, A	328	27	58.7	341	11	US-11-188-298-8905	Sequence 8905, Ap
256	27	58.7	324	9	US-10-714-887-300	Sequence 300, App	329	27	58.7	341	11	US-11-188-298-8949	Sequence 8949, Ap
257	27	58.7	326	11	US-11-188-298-4969	Sequence 4969, Ap	330	27	58.7	342	11	US-11-188-298-19513	Sequence 19513, A
258	27	58.7	326	11	US-11-199-233-8	Sequence 8, Appl	331	27	58.7	342	11	US-11-188-298-15139	Sequence 15139, A
259	27	58.7	328	11	US-11-087-099-4126	Sequence 4126, Ap	332	27	58.7	342	11	US-11-188-298-10269	Sequence 10269, A
260	27	58.7	331	11	US-11-188-298-3201	Sequence 3201, Ap	333	27	58.7	342	11	US-11-188-298-18496	Sequence 18496, A
261	27	58.7	331	11	US-11-188-298-3228	Sequence 3228, Ap	334	27	58.7	342	11	US-11-188-298-18496	Sequence 18496, A
262	27	58.7	332	11	US-11-183-664-1	Sequence 1, Appl	335	27	58.7	343	11	US-11-188-298-2561	Sequence 2561, Ap
263	27	58.7	332	11	US-11-188-298-551	Sequence 551, App	336	27	58.7	345	11	US-11-188-298-16881	Sequence 16881, A
264	27	58.7	332	11	US-11-188-298-13876	Sequence 13876, A	337	27	58.7	345	11	US-11-188-298-21305	Sequence 21305, A
265	27	58.7	332	11	US-11-188-298-21933	Sequence 21933, A	338	27	58.7	347	11	US-11-188-298-12383	Sequence 12383, A
266	27	58.7	333	11	US-11-183-664-6	Sequence 6, Appl	339	27	58.7	350	11	US-11-188-298-12989	Sequence 12989, A
267	27	58.7	334	11	US-11-140-417-8	Sequence 8, Appl	340	27	58.7	352	11	US-11-188-298-20	Sequence 20, Appl
268	27	58.7	335	8	US-10-511-937-2536	Sequence 2536, App	341	27	58.7	353	11	US-11-188-298-18005	Sequence 18005, A
269	27	58.7	335	9	US-10-511-937-2560	Sequence 2560, App	342	27	58.7	354	9	US-10-838-616-16	Sequence 16, Appl
270	27	58.7	335	8	US-10-995-561-704	Sequence 704, App	343	27	58.7	357	11	US-11-188-298-10253	Sequence 10253, A
271	27	58.7	335	11	US-11-141-947-2	Sequence 2, Appl	344	27	58.7	360	11	US-11-188-298-12322	Sequence 12322, A
272	27	58.7	337	11	US-11-188-298-717	Sequence 717, App	345	27	58.7	363	9	US-10-838-616-10	Sequence 10, Appl
273	27	58.7	337	11	US-11-188-298-846	Sequence 846, App	346	27	58.7	365	11	US-11-188-298-11126	Sequence 11126, A
274	27	58.7	337	11	US-11-188-298-1785	Sequence 1785, App	347	27	58.7	366	11	US-11-098-666-11126	Sequence 11126, A
275	27	58.7	337	11	US-11-188-298-2392	Sequence 2392, Ap	348	27	58.7	370	11	US-11-188-298-14660	Sequence 14660, A
276	27	58.7	337	11	US-11-188-298-2558	Sequence 2558, Ap	349	27	58.7	385	11	US-11-188-298-3041	Sequence 3041, Ap
277	27	58.7	337	11	US-11-188-298-2570	Sequence 2570, Ap	350	27	58.7	385	11	US-11-188-298-10855	Sequence 10855, A
278	27	58.7	337	11	US-11-188-298-2664	Sequence 2664, Ap	351	27	58.7	386	11	US-11-188-298-14049	Sequence 14049, A
279	27	58.7	337	11	US-11-188-298-2954	Sequence 2954, Ap	352	27	58.7	386	11	US-11-188-298-20493	Sequence 20493, A
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284	27	58.7	337	11	US-11-188-298-7167	Sequence 7167, Ap	357	27	58.7	406	11	US-11-188-298-8140	Sequence 8140, Ap
285	27	58.7	337	11	US-11-188-298-7573	Sequence 7573, Ap	358	27	58.7	406	11	US-11-188-298-15109	Sequence 15109, A
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287	27	58.7	337	11	US-11-188-298-8680	Sequence 8680, Ap	360	27	58.7	415	11	US-11-079-463-8796	Sequence 8797, Ap
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313	27	58.7	338	11	US-11-188-298-5280	Sequence 5280, Ap							

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389	27	58.7	631	11	US-11-188-298-865	Sequence 865, App	462	26	56.5	257	11	US-11-188-298-11558	Sequence 21538, A
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391	27	58.7	647	11	US-11-079-463-6814	Sequence 6814, Ap	464	26	56.5	260	9	US-10-497-135-24	Sequence 24, App1
392	27	58.7	679	11	US-11-079-463-7773	Sequence 7773, Ap	465	26	56.5	260	11	US-11-188-298-6417	Sequence 6417, Ap
393	27	58.7	713	8	US-10-005-928-345	Sequence 345, App	466	26	56.5	260	11	US-11-269-215-23	Sequence 21, App1
394	27	58.7	781	11	US-11-072-512-2937	Sequence 2937, Ap	467	26	56.5	260	11	US-11-269-215-24	Sequence 22, App1
395	27	58.7	781	11	US-11-188-298-5388	Sequence 5388, Ap	468	26	56.5	261	11	US-11-188-298-6910	Sequence 6910, Ap
396	27	58.7	813	9	US-10-877-346-45	Sequence 45, App1	469	26	56.5	261	11	US-11-005-004-2092	Sequence 2240, Ap
397	27	58.7	819	11	US-11-045-004-778	Sequence 778, App	470	26	56.5	262	11	US-11-188-298-12240	Sequence 1240, Ap
398	27	58.7	820	11	US-11-147-047-31	Sequence 31, App1	471	26	56.5	263	11	US-11-188-298-9786	Sequence 9786, Ap
399	27	58.7	860	11	US-11-019-711-59	Sequence 59, App1	472	26	56.5	267	11	US-11-188-298-15650	Sequence 15650, A
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404	27	58.7	931	11	US-11-019-711-119	Sequence 119, App	477	26	56.5	273	11	US-11-188-298-1754	Sequence 1754, Ap
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433	26	56.5	142	9	US-10-467-657-1558	Sequence 1558, Ap	506	26	56.5	296	9	US-10-714-887-226	Sequence 226, App
434	26	56.5	149	9	US-10-530-253-18	Sequence 18, App1	507	26	56.5	299	11	US-11-188-298-51902	Sequence 51902, Ap
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554	26	56.5	319	9	US-10-454-437-116	Sequence 116, App
555	26	56.5	319	11	US-11-188-298-20507	Sequence 20507, A
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863	25	54.3	194	11	US-11-188-298-9187	Sequence 9187, App	936	25	54.3	221	11	US-11-096-568A-1564	Sequence 3581, App
864	25	54.3	195	11	US-11-087-099-6517	Sequence 6517, App	937	25	54.3	225	11	US-11-087-099-3581	Sequence 1037, App
865	25	54.3	195	11	US-11-087-099-8980	Sequence 8980, App	938	25	54.3	225	11	US-11-096-568A-1037	Sequence 14358, A
866	25	54.3	195	11	US-11-188-298-6938	Sequence 6938, App	939	25	54.3	226	11	US-11-188-298-14358	Sequence 60, App1
867	25	54.3	195	11	US-11-188-298-8407	Sequence 8407, App	940	25	54.3	226	11	US-10-703-1999-60	Sequence 23896, A
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870	25	54.3	196	11	US-11-188-298-1164	Sequence 1164, App	943	25	54.3	226	11	US-11-096-568A-24646	Sequence 6657, App
871	25	54.3	196	11	US-11-188-298-3931	Sequence 3931, App	944	25	54.3	226	11	US-11-188-298-11812	Sequence 11812, A
872	25	54.3	196	11	US-11-188-298-4987	Sequence 4987, App	945	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
873	25	54.3	196	11	US-11-188-298-5360	Sequence 5360, App	946	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
874	25	54.3	196	11	US-11-188-298-6399	Sequence 6399, App	947	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
875	25	54.3	196	11	US-11-188-298-7779	Sequence 7779, App	948	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
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879	25	54.3	196	11	US-11-188-298-9398	Sequence 9398, App	952	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
880	25	54.3	196	11	US-11-188-298-9473	Sequence 9473, App	953	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
881	25	54.3	196	11	US-11-188-298-9825	Sequence 9825, App	954	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
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883	25	54.3	196	11	US-11-188-298-11734	Sequence 11734, App	956	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
884	25	54.3	196	11	US-11-188-298-11998	Sequence 11998, App	957	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
885	25	54.3	196	11	US-11-188-298-13537	Sequence 13537, App	958	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
886	25	54.3	196	11	US-11-188-298-14488	Sequence 14488, App	959	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
887	25	54.3	196	11	US-11-188-298-14781	Sequence 14781, App	960	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
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889	25	54.3	196	11	US-11-188-298-16427	Sequence 16427, App	962	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
890	25	54.3	196	11	US-11-188-298-16859	Sequence 16859, App	963	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
891	25	54.3	196	11	US-11-188-298-17886	Sequence 17886, App	964	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
892	25	54.3	196	11	US-11-188-298-18851	Sequence 18851, App	965	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
893	25	54.3	196	11	US-11-188-298-20944	Sequence 20944, App	966	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
894	25	54.3	196	11	US-11-188-298-21196	Sequence 21196, App	967	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
895	25	54.3	196	11	US-11-188-298-21772	Sequence 21772, App	968	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A
896	25	54.3	196	11	US-11-188-298-21821	Sequence 21821, App	969	25	54.3	226	11	US-11-096-568A-6657	Sequence 6657, App
897	25	54.3	196	11	US-11-188-298-22141	Sequence 22141, App	970	25	54.3	226	11	US-11-096-568A-6657	Sequence 11812, A

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911 25 54.3 254 9 US-10-995-561-841 Sequence 841, App
912 25 54.3 256 9 US-10-537-002-64 Sequence 64, App
913 25 54.3 256 9 US-10-194-487-374 Sequence 374, App
914 25 54.3 256 9 US-10-195-883-374 Sequence 374, App
915 25 54.3 256 9 US-10-195-883-374 Sequence 374, App
916 25 54.3 256 9 US-10-195-883-374 Sequence 374, App
917 25 54.3 256 11 US-11-096-568A-24913 Sequence 24913, A
918 25 54.3 258 9 US-10-793-626-170 Sequence 170, App
919 25 54.3 258 9 US-10-793-626-236 Sequence 236, App
920 25 54.3 258 9 US-10-793-626-1614 Sequence 1614, App
921 25 54.3 260 11 US-11-079-463-9916 Sequence 9916, App
922 25 54.3 260 11 US-11-079-463-9916 Sequence 9916, App
923 25 54.3 264 11 US-11-096-568A-21268 Sequence 21268, A
924 25 54.3 264 11 US-11-188-298-1485 Sequence 1485, App
925 25 54.3 267 11 US-11-087-099-10418 Sequence 10418, A
926 25 54.3 267 11 US-11-087-099-10418 Sequence 10418, A
927 25 54.3 268 11 US-11-188-298-21091 Sequence 21091, A
928 25 54.3 269 11 US-11-045-004-479 Sequence 479, App
929 25 54.3 270 11 US-11-096-568A-4898 Sequence 4898, App
930 25 54.3 271 11 US-11-096-568A-8711 Sequence 8711, App
931 25 54.3 272 11 US-11-087-099-8189 Sequence 8189, App
932 25 54.3 273 11 US-11-188-298-22010 Sequence 22010, A
933 25 54.3 275 9 US-10-821-234-1013 Sequence 1013, App
934 25 54.3 276 11 US-11-087-099-6624 Sequence 6624, App
935 25 54.3 276 11 US-11-087-099-6624 Sequence 6624, App
936 25 54.3 276 11 US-11-087-099-5099 Sequence 5099, App
937 25 54.3 277 11 US-11-188-298-3428 Sequence 3428, App
938 25 54.3 277 11 US-11-188-298-5318 Sequence 5318, App
939 25 54.3 277 11 US-11-188-298-6156 Sequence 6156, App
940 25 54.3 277 11 US-11-188-298-6156 Sequence 6156, App
1000 25 54.3 277 11 US-11-188-298-14294 Sequence 14294, A
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ALIGNMENTS

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RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; TYPE: PRT
; LENGTH: 151
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13

Query Match 100.0%; Score 46; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.032;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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; GENERAL INFORMATION:
; APPLICANT: Healthbanc Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; PRIOR FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; TYPE: PRT
; LENGTH: 158
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3

Query Match 100.0%; Score 46; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.034;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-1

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
US-10-530-253-3

Query Match 100.0%; Score 46; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.032;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
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| | | | |
Db 18 ELQTTTHDI 26

RESULT 5
US-10-530-253-5
; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
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| | | | |
Db 18 ELQTTTHDI 26

RESULT 6
US-10-530-253-7
; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 248

;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
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| | | | |
Db 115 ELQTTTHDI 123

RESULT 7
US-10-530-253-9
; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 46; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 ELQTTTHDI 9
| | | | |
| | | | |
Db 115 ELQTTTHDI 123

RESULT 8
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 46; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHD 9
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Db 115 ELQTTIHD 123

RESULT 9

US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XINOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; CURRENT FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: Patentin Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 46; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.057;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHD 9
|||
Db 123 ELQTTIHD 131

RESULT 10

US-11-096-568A-1943
; Sequence 1943, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1943
; LENGTH: 250
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(250)
; OTHER INFORMATION: Ceres Seq. ID no. 15180148
US-11-096-568A-1943

Query Match 87.0%; Score 40; DB 11; Length 250;
Best Local Similarity 87.5%; Pred. No. 0.96;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHD 8
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Db 180 ELQTTIHD 187

RESULT 11

US-11-096-568A-1942

; Sequence 1942, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1942
; LENGTH: 298
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(298)
; OTHER INFORMATION: Ceres Seq. ID no. 15180147
US-11-096-568A-1942

Query Match 87.0%; Score 40; DB 11; Length 298;
Best Local Similarity 87.5%; Pred. No. 1.2;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHD 8
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Db 228 ELQTTIHD 235

RESULT 12

US-11-096-568A-1941
; Sequence 1941, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1941
; LENGTH: 303
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(303)
; OTHER INFORMATION: Ceres Seq. ID no. 15180146
US-11-096-568A-1941

Query Match 87.0%; Score 40; DB 11; Length 303;
Best Local Similarity 87.5%; Pred. No. 1.2;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 ELQTTIHD 8
|||
Db 233 ELQTTIHD 240

RESULT 13

US-10-503-575-174
; Sequence 174, Application US/10503575
; Publication No. US20050244823A1
; GENERAL INFORMATION:
; APPLICANT: Drifhout, Jan Wouter
; APPLICANT: van Veele, Petrus Antonius
; APPLICANT: Koning, Frits
; TITLE OF INVENTION: NOVEL EPITOPES FOR CELIAC DISEASE AND AUTOIMMUNE DISEASES, METHOD
; TITLE OF INVENTION: DETECTING THOSE AND NOVEL NON-ANTIGENIC FOOD COMPOUNDS
; FILE REFERENCE: 2799/72843-PCT-US
; CURRENT APPLICATION NUMBER: US/10/503,575
; CURRENT FILING DATE: 2004-08-04

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; PRIOR APPLICATION NUMBER: PCT/NL03/00077
; PRIOR FILING DATE: 2003-02-04
; PRIOR APPLICATION NUMBER: EP 02075456.0
; PRIOR FILING DATE: 2002-02-04
; NUMBER OF SEQ ID NOS: 340
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 174
; LENGTH: 19
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-503-575-174

Query Match      76.1%; Score 35; DB 9; Length 19;
Best Local Similarity 75.0%; Pred. No. 0.53;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 ELQTTIHD 8
Db      8 ELQATLHD 15

RESULT 14
US-10-986-405-199
; Sequence 199, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; PRIOR FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 199
; LENGTH: 133
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-199

Query Match      76.1%; Score 35; DB 9; Length 133;
Best Local Similarity 75.0%; Pred. No. 4.9;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 ELQTTIHD 8
Db      106 ELQATLHD 113

RESULT 15
US-10-986-405-205
; Sequence 205, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
```

```

; CURRENT FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 205
; LENGTH: 135
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-205

Query Match      76.1%; Score 35; DB 9; Length 135;
Best Local Similarity 75.0%; Pred. No. 5;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 ELQTTIHD 8
Db      106 ELQATLHD 113

RESULT 16
US-10-986-405-229
; Sequence 229, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; PRIOR FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 229
; LENGTH: 140
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-229

Query Match      76.1%; Score 35; DB 9; Length 140;
Best Local Similarity 75.0%; Pred. No. 5.2;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 ELQTTIHD 8
```

Db 121 ELQATLHD 128

RESULT 17

US-10-986-405-200
; Sequence 200, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; PRIOR FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 200
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (50)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (122)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (138)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (143)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-10-986-405-200

Query Match 76.1%; Score 35; DB 9; Length 149;
Best Local Similarity 75.0%; Pred. No. 5.6;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
Db 106 ELQATLHD 113

RESULT 18
US-10-986-405-216
; Sequence 216, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; CURRENT FILING DATE: 2004-11-12

; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 216
; LENGTH: 184
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (9)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (50)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (98)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (137)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
; FEATURE:
; NAME/KEY: SITE
; LOCATION: (163)
; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-10-986-405-216

Query Match 76.1%; Score 35; DB 9; Length 184;
Best Local Similarity 75.0%; Pred. No. 7.1;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHD 8
Db 106 ELQATLHD 113

RESULT 19
US-11-096-568A-19577
; Sequence 19577, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19577
; LENGTH: 287
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(287)
; OTHER INFORMATION: Ceres Seq. ID no. 12373656

US-11-096-568A-19577

Query Match 73.9%; Score 34; DB 11; Length 287;
Best Local Similarity 75.0%; Pred. No. 19;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
: : : : :
Db 94 VKTTIHDI 101

RESULT 20

US-10-703-799B-62
; Sequence 62, Application US/10703799B
; Publication No. US20060078884A1
; GENERAL INFORMATION:
; APPLICANT: Pompejus, Markus
; APPLICANT: Kroger, Burkhard
; APPLICANT: Schroder, Hartwig
; APPLICANT: Zeider, Oskar
; APPLICANT: Haberauer, Gregor
; APPLICANT: Lee, Heung-Shick
; APPLICANT: Kim, Hyung-Joon
; TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING STRESS,
; FILE REFERENCE: BGI-124CPCN
; CURRENT APPLICATION NUMBER: US/10/703,799B
; PRIOR FILING DATE: 2003-11-07
; PRIOR APPLICATION NUMBER: 09/603,208
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: 60/141031
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: 60/142692
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: 60/151214
; PRIOR FILING DATE: 1999-08-27
; PRIOR APPLICATION NUMBER: DE 19930429.7
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: DE 19931413.6
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19931457.8
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19931541.8
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19932209.0
; PRIOR FILING DATE: 1999-07-09
; PRIOR APPLICATION NUMBER: DE 19932230.9
; PRIOR FILING DATE: 1999-07-09
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 306
; SEQ ID NO 62
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
; US-10-703-799B-62

Query Match 73.9%; Score 34; DB 9; Length 383;
Best Local Similarity 66.7%; Pred. No. 26;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTIHDI 9
: : : : :
Db 237 EMKTIHDI 245

RESULT 21

US-11-096-568A-19576
; Sequence 19576, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Thedy

FILE REFERENCE: 2750-1592PUS2

; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19576
; LENGTH: 448

; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(448)
; OTHER INFORMATION: Ceres Seq. ID no. 12373655
US-11-096-568A-19576

Query Match 73.9%; Score 34; DB 11; Length 448;
Best Local Similarity 75.0%; Pred. No. 31;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
: : : : :
Db 255 VKTTIHDI 262

RESULT 22

US-11-096-568A-19575
; Sequence 19575, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19575
; LENGTH: 463
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(463)
; OTHER INFORMATION: Ceres Seq. ID no. 12373654
US-11-096-568A-19575

Query Match 73.9%; Score 34; DB 11; Length 463;
Best Local Similarity 75.0%; Pred. No. 33;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTTIHDI 9
: : : : :
Db 270 VKTTIHDI 277

RESULT 23

US-11-080-991-108
; Sequence 108, Application US/11080991
; Publication No. US20050266437A1
; GENERAL INFORMATION:
; APPLICANT: Velby, Petter Ole
; TITLE OF INVENTION: COMPOSITIONS, KITS, AND METHODS FOR
; TITLE OF INVENTION: IDENTIFICATION, ASSESSMENT, PREVENTION, AND THERAPY OF BREAST
; FILE REFERENCE: MRI-039
; CURRENT APPLICATION NUMBER: US/11/080,991
; CURRENT FILING DATE: 2005-03-11
; PRIOR APPLICATION NUMBER: US/10/176,847
; PRIOR FILING DATE: 2002-06-21
; NUMBER OF SEQ ID NOS: 112
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 108
; LENGTH: 628
; TYPE: PRT

```
; ORGANISM: Homo sapiens
US-11-080-991-108

Query Match
Best Local Similarity 73.9%; Score 34; DB 11; Length 628;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

OY 1 ELQTTIHD 8
DB 87 ELQVTIHD 94

RESULT 24
US-10-204-639-4
; Sequence 4, Application US/10204639
; Publication No. US20060063152A1
; GENERAL INFORMATION:
; APPLICANT: Ogamu Ohara
; APPLICANT: Takahiro Nagase
; APPLICANT: Daijuke Nakajima
; TITLE OF INVENTION: NOVEL GENE AND PROTEIN ENCODED BY THE GENE
; FILE REFERENCE: PH-1416-PCT
; CURRENT APPLICATION NUMBER: US/10/204,639
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: JP 2000-389742
; PRIOR FILING DATE: 2000-12-22
; PRIOR APPLICATION NUMBER: JP 2001-095524
; PRIOR FILING DATE: 2001-03-29
; PRIOR APPLICATION NUMBER: JP 2001-127066
; PRIOR FILING DATE: 2001-04-25
; NUMBER OF SEQ ID NOS: 140
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 4
; LENGTH: 1479
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-204-639-4

Query Match
Best Local Similarity 73.9%; Score 34; DB 9; Length 1479;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 ELQTTIHD 9
DB 1003 ELQTTIHD 1011

RESULT 25
US-10-877-346-13
; Sequence 13, Application US/10877346
; Publication No. US20060014153A1
; GENERAL INFORMATION:
; APPLICANT: Gerlach, Valerie L
; APPLICANT: MacDougall, John R
; APPLICANT: Smithson, Glenda
; APPLICANT: Millet, Isabelle
; APPLICANT: Stone, David
; APPLICANT: Gunther, Erik
; APPLICANT: Ellerman, Karen
; APPLICANT: Grosse, William M
; APPLICANT: Alsobrook II, John P
; APPLICANT: Lepley, Denise M
; APPLICANT: Burgess, Catherine E
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Kekuda, Ramesh
; APPLICANT: Spytek, Kimberly A
; APPLICANT: Leach, Martin D
; APPLICANT: Shinkere, Richard A
; TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-124
; CURRENT APPLICATION NUMBER: US/10/877,346
; PRIOR FILING DATE: 2004-06-25
; PRIOR APPLICATION NUMBER: US/09/964,956
```

```
; PRIOR FILING DATE: 2001-09-26
; PRIOR APPLICATION NUMBER: 60/235,631
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/235,633
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/235,808
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,064
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,065
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,066
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,135
; PRIOR FILING DATE: 2000-09-28
; PRIOR APPLICATION NUMBER: 60/237,434
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/238,321
; PRIOR FILING DATE: 2000-10-05
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 127
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 13
; LENGTH: 1896
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-877-346-13

Query Match
Best Local Similarity 73.9%; Score 34; DB 9; Length 1896;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 ELQTTIHD 9
DB 1295 ELQTTIHD 1303

RESULT 26
US-10-877-346-44
; Sequence 44, Application US/10877346
; Publication No. US20060014153A1
; GENERAL INFORMATION:
; APPLICANT: Gerlach, Valerie L
; APPLICANT: MacDougall, John R
; APPLICANT: Smithson, Glenda
; APPLICANT: Millet, Isabelle
; APPLICANT: Stone, David
; APPLICANT: Gunther, Erik
; APPLICANT: Ellerman, Karen
; APPLICANT: Grosse, William M
; APPLICANT: Alsobrook II, John P
; APPLICANT: Lepley, Denise M
; APPLICANT: Burgess, Catherine E
; APPLICANT: Padigaru, Muralidhara
; APPLICANT: Kekuda, Ramesh
; APPLICANT: Spytek, Kimberly A
; APPLICANT: Leach, Martin D
; APPLICANT: Shinkere, Richard A
; TITLE OF INVENTION: Novel Proteins and Nucleic Acids Encoding Same
; FILE REFERENCE: 21402-124
; CURRENT APPLICATION NUMBER: US/10/877,346
; PRIOR FILING DATE: 2004-06-25
; PRIOR APPLICATION NUMBER: US/09/964,956
; PRIOR FILING DATE: 2001-09-26
; PRIOR APPLICATION NUMBER: 60/235,631
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/235,633
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/235,808
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,064
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,065
```



```

; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,066
; PRIOR FILING DATE: 2000-09-27
; PRIOR APPLICATION NUMBER: 60/236,135
; PRIOR FILING DATE: 2000-09-28
; PRIOR APPLICATION NUMBER: 60/237,434
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/238,321
; PRIOR FILING DATE: 2000-10-05
; PRIOR APPLICATION NUMBER: 60/238,321
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 127
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 44
; LENGTH: 1905
; TYPE: PRT
; ORGANISM: Xenopus laevis
US-10-877-346-44
```

```
Query Match      73.9%; Score 34; DB 9; Length 1905;
Best Local Similarity 66.7%; Pred. No. 1.6e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LQTTIHDI 9
      : |||:|:
Db      1292 ELQTDIHEL 1300
```

```

RESULT 27
US-10-530-253-17
; Sequence 17, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 17
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 33
US-10-530-253-17
```

```
Query Match      71.7%; Score 33; DB 9; Length 149;
Best Local Similarity 75.0%; Pred. No. 14;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 LQTTIHDI 9
      : |||:|:
Db      19 LQTTIHNI 26
```

```

RESULT 28
US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
```

```

; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 26
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 68
US-10-530-253-26
```

```
Query Match      71.7%; Score 33; DB 9; Length 158;
Best Local Similarity 62.5%; Pred. No. 15;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      2 LQTTIHDI 9
      : |||:|:
Db      21 LQTTIHDI 28
```

```

RESULT 29
US-10-506-454-1603
; Sequence 1603, Application US/10506454
; Publication No. US20060068386A1
; GENERAL INFORMATION:
; APPLICANT: Slesarev, Alexi I
; APPLICANT: Mezheva, Katja V
; APPLICANT: Polushin, Nikolai N
; APPLICANT: Shcherbina, Olga V
; APPLICANT: Shakhova, Vera V
; APPLICANT: Malykh, Andrei G
; APPLICANT: Kozavkin, Sergei A
; TITLE OF INVENTION: The Complete Genome and Protein Sequences of the Hyperthermophilic
; TITLE OF INVENTION: Methanopyrus Kandleri AV19 and Monophyly of Archaeal Methanogens
; FILE REFERENCE: FID001
; CURRENT APPLICATION NUMBER: US/10/506,454
; PRIOR FILING DATE: 2004-08-31
; PRIOR APPLICATION NUMBER: PCT/US03/06664
; PRIOR FILING DATE: 2003-03-04
; PRIOR APPLICATION NUMBER: 60/361,742
; PRIOR FILING DATE: 2002-03-04
; NUMBER OF SEQ ID NOS: 1722
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1603
; LENGTH: 237
; TYPE: PRT
; ORGANISM: Methanopyrus kandleri
US-10-506-454-1603
```

```
Query Match      71.7%; Score 33; DB 9; Length 237;
Best Local Similarity 62.5%; Pred. No. 24;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      2 LQTTIHDI 9
      : |||:|:
Db      167 IHTTVHDI 174
```

```

RESULT 30
US-10-504-120-21
; Sequence 21, Application US/10504120
; Publication No. US20060088829A1
; GENERAL INFORMATION:
; APPLICANT: Exelixis, Inc.
; TITLE OF INVENTION: MINRA AS MODIFIERS OF INSULIN RECEPTOR SIGNALING AND METHODS OF
; FILE REFERENCE: EX03-003C-PC
; CURRENT APPLICATION NUMBER: US/10/504,120
; CURRENT FILING DATE: 2004-08-06
; PRIOR APPLICATION NUMBER: 60/354,824
```

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; PRIOR FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 60/358,217
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,189
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,126
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,995
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/358,756
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/358,765
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/359,531
; PRIOR FILING DATE: 2002-02-25
; PRIOR APPLICATION NUMBER: 60/360,222
; PRIOR FILING DATE: 2002-02-26
; PRIOR APPLICATION NUMBER: 60/360,224
; PRIOR FILING DATE: 2002-02-26
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 21
; LENGTH: 1763
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-504-120-21
```

```
Query Match          71.7%; Score 33; DB 8; Length 1763;
Best Local Similarity 55.6%; Pred. No. 2.4e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
```

```
QY 1 ELQTTIHDI 9
DB 379 ELRTIVHDL 387
```

```
RESULT 31
US-10-504-120-22
; Sequence 22, Application US/10504120
; Publication No. US2006008829A1
; GENERAL INFORMATION:
; APPLICANT: Exeligen, Inc.
; TITLE OF INVENTION: MINIS AS MODIFIERS OF INSULIN RECEPTOR SIGNALING AND METHODS OF
; FILE REFERENCE: EX03-003C-PC
; CURRENT APPLICATION NUMBER: US/10/504,120
; CURRENT FILING DATE: 2004-08-06
; PRIOR APPLICATION NUMBER: 60/354,824
; PRIOR FILING DATE: 2002-02-06
; PRIOR APPLICATION NUMBER: 60/358,217
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,189
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,126
; PRIOR FILING DATE: 2002-02-20
; PRIOR APPLICATION NUMBER: 60/358,995
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/358,756
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/358,765
; PRIOR FILING DATE: 2002-02-21
; PRIOR APPLICATION NUMBER: 60/359,531
; PRIOR FILING DATE: 2002-02-25
; PRIOR APPLICATION NUMBER: 60/360,222
; PRIOR FILING DATE: 2002-02-26
; PRIOR APPLICATION NUMBER: 60/360,224
; PRIOR FILING DATE: 2002-02-26
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 32
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 22
; LENGTH: 1807
```

```
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-504-120-22
```

```
Query Match          71.7%; Score 33; DB 8; Length 1807;
Best Local Similarity 55.6%; Pred. No. 2.5e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
```

```
QY 1 ELQTTIHDI 9
DB 379 ELRTIVHDL 387
```

```
RESULT 32
US-11-087-099-2617
; Sequence 2617, Application US/11087099
; Publication No. US2006004196A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 2617
; LENGTH: 162
; TYPE: PRT
; ORGANISM: Corynebacterium efficiens YS-314
US-11-087-099-2617
```

```
Query Match          69.6%; Score 32; DB 11; Length 162;
Best Local Similarity 66.7%; Pred. No. 25;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
QY 1 ELQTTIHDI 9
DB 2 EAMTSIHDI 10
```

```
RESULT 33
US-11-188-298-17532
; Sequence 17532, Application US/11188298
; Publication No. US2006007552A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 17532
; LENGTH: 333
; TYPE: PRT
; ORGANISM: Magnetococcus sp. MC-1
US-11-188-298-17532
```

```
Query Match          69.6%; Score 32; DB 11; Length 333;
Best Local Similarity 100.0%; Pred. No. 57;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 4 TTTHDI 9
DB 175 TTTHDI 180
```

```
RESULT 34
US-11-188-298-16137
; Sequence 16137, Application US/11188298
; Publication No. US2006007552A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
```

FILE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 16137
LENGTH: 337
TYPE: PRT
ORGANISM: *Shewanella oneidensis* MR-1
US-11-188-298-16137

Query Match 69.6%; Score 32; DB 11; Length 337;
Best Local Similarity 100.0%; Pred. No. 58;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTDDI 9
DB 176 TTTTDDI 181

RESULT 35
US-11-188-298-6731
Sequence 6731, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 6731
LENGTH: 340
TYPE: PRT
ORGANISM: *SYNECHOCOCCUS* SP. MH 8102
US-11-188-298-6731

Query Match 69.6%; Score 32; DB 11; Length 340;
Best Local Similarity 100.0%; Pred. No. 59;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTDDI 9
DB 181 TTTTDDI 186

RESULT 36
US-11-188-298-3477
Sequence 3477, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 3477
LENGTH: 498
TYPE: PRT
ORGANISM: *Nicotiana tabacum*
US-11-188-298-3477

Query Match 69.6%; Score 32; DB 11; Length 498;
Best Local Similarity 55.6%; Pred. No. 91;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 ELQTTTDDI 9
DB 152 ELQSLDHL 160

RESULT 37
US-10-873-528-37
Sequence 37, Application US/10873528
Publication No. US20050276814A1
GENERAL INFORMATION:
APPLICANT: Microbial Technics Limited
APPLICANT: Gilbert, Christophe FG
APPLICANT: Hansbro, Philip M
TITLE OF INVENTION: Proteins
FILE REFERENCE: PWC/P21129MO
CURRENT APPLICATION NUMBER: US/10/873,528
CURRENT FILING DATE: 2004-06-23
PRIOR APPLICATION NUMBER: US/09/769,787
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: GB 9816337.1
PRIOR FILING DATE: 1998-03-27
PRIOR APPLICATION NUMBER: US 60/125164
PRIOR FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 388
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 37
LENGTH: 513
TYPE: PRT
ORGANISM: *Streptococcus pneumoniae*
US-10-873-528-37

Query Match 69.6%; Score 32; DB 9; Length 513;
Best Local Similarity 100.0%; Pred. No. 94;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTDDI 9
DB 495 TTTTDDI 500

RESULT 38
US-10-873-528-193
Sequence 193, Application US/10873528
Publication No. US20050276814A1
GENERAL INFORMATION:
APPLICANT: Microbial Technics Limited
APPLICANT: Gilbert, Christophe FG
APPLICANT: Hansbro, Philip M
TITLE OF INVENTION: Proteins
FILE REFERENCE: PWC/P21129MO
CURRENT APPLICATION NUMBER: US/10/873,528
CURRENT FILING DATE: 2004-06-23
PRIOR APPLICATION NUMBER: US/09/769,787
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: GB 9816337.1
PRIOR FILING DATE: 1998-03-27
PRIOR APPLICATION NUMBER: US 60/125164
PRIOR FILING DATE: 1999-03-19
NUMBER OF SEQ ID NOS: 388
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 193
LENGTH: 513
TYPE: PRT
ORGANISM: *Streptococcus pneumoniae*
US-10-873-528-193

Query Match 69.6%; Score 32; DB 9; Length 513;
Best Local Similarity 100.0%; Pred. No. 94;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTDDI 9
DB 495 TTTTDDI 500

```

RESULT 39
US-11-103-957-61
; Sequence 61, Application US/11103957
; Publication No. US20050281847A1
; GENERAL INFORMATION:
; APPLICANT: Berthet, Francois-Xavier Jacques
; APPLICANT: Lobet, Yves
; APPLICANT: Poolman, Jan
; APPLICANT: Verlant, Vincent Georges Christian Louis
; TITLE OF INVENTION: Vaccine Composition
; FILE REFERENCE: B45261
; CURRENT APPLICATION NUMBER: US/11/103,957
; CURRENT FILING DATE: 2005-04-12
; PRIOR APPLICATION NUMBER: US/10/467,534
; PRIOR FILING DATE: 2004-02-03
; PRIOR APPLICATION NUMBER: PCT/EP02/01356
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: GB 0103169.9
; PRIOR FILING DATE: 2001-02-08
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 61
; LENGTH: 553
; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-11-103-957-61

Query Match
Best Local Similarity 69.6%; Score 32; DB 11; Length 553;
Pred. No. 1e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 LQTHDI 9
|:|||||
Db 284 LETLHDI 291

RESULT 40
US-11-087-099-1991
; Sequence 1991, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 1991
; LENGTH: 643
; TYPE: PRT
; ORGANISM: Magnetospirillum magnetotacticum
US-11-087-099-1991

Query Match
Best Local Similarity 69.6%; Score 32; DB 11; Length 643;
Pred. No. 1.2e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 LQTHDI 8
|:|||||
Db 5 LQTHDI 11

RESULT 41
US-11-188-298-5563
; Sequence 5563, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298

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; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 5563
; LENGTH: 989
; TYPE: PRT
; ORGANISM: Trichodesmium erythraeum IMS101
; FEATURES:
; NAME/KEY: unsure
; LOCATION: (1)..(989)
; OTHER INFORMATION: unsure at all Xaa locations
US-11-188-298-5563

Query Match
Best Local Similarity 69.6%; Score 32; DB 11; Length 989;
Pred. No. 2e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 3 QTHHDI 9
|:|||||
Db 457 KTHHDI 463

RESULT 42
US-11-096-568A-6040
; Sequence 6040, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6040
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURES:
; NAME/KEY: misc_feature
; LOCATION: (1)..(329)
; OTHER INFORMATION: Ceres Seq. ID no. 14313169
US-11-096-568A-6040

Query Match
Best Local Similarity 67.4%; Score 31; DB 11; Length 329;
Pred. No. 90;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 LQTHDI 8
|:|||||
Db 203 LQTHDI 209

RESULT 43
US-11-188-298-21765
; Sequence 21765, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 21765
; LENGTH: 335
; TYPE: PRT
; ORGANISM: Rhodobacter sphaeroides
US-11-188-298-21765

```

Query Match 67.4%; Score 31; DB 11; Length 335;
Best Local Similarity 83.3%; Pred. No. 92;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 4 TTTTTHDI 9
|||
DB 176 TTTTHDV 181

RESULT 44

US-11-188-298-16409
; Sequence 16409, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16409
; LENGTH: 340
; TYPE: PRT
; ORGANISM: *Prochlorococcus marinus* subsp. *paetoris* str. CCMP1378
US-11-188-298-16409

Query Match 67.4%; Score 31; DB 11; Length 340;
Best Local Similarity 83.3%; Pred. No. 94;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 4 TTTTTHDI 9
|||
DB 181 TTTTHDV 186

RESULT 45

US-11-096-568A-6039
; Sequence 6039, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6039
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURES:
; NAME/KEY: misc feature
; LOCATION: (1)...(540)
; OTHER INFORMATION: Ceres Seq. ID no. 14313168
US-11-096-568A-6039

Query Match 67.4%; Score 31; DB 11; Length 540;
Best Local Similarity 71.4%; Pred. No. 1.6e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

OY 2 LOTTTHDI 8
|||
DB 414 LOTSLSHD 420

RESULT 46
US-11-169-041-145
; Sequence 145, Application US/11169041
; Publication No. US20060019284A1
; GENERAL INFORMATION:

APPLICANT: Bristol-Myers Squibb Company
; TITLE OF INVENTION: IDENTIFICATION OF POLYNUCLEOTIDES FOR PREDICTING ACTIVITY OF
; TITLE OF INVENTION: COMPOUNDS THAT INTERACT WITH AND/OR MODULATE PROTEIN TYROSINE
; TITLE OF INVENTION: KINASES AND/OR PROTEIN TYROSINE KINASE PATHWAYS IN LUNG CANCER
; FILE REFERENCE: 10001 NP
; CURRENT APPLICATION NUMBER: US/11/169,041
; CURRENT FILING DATE: 2005-06-28
; PRIOR APPLICATION NUMBER: 60/584,405
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 527
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 145
; LENGTH: 917
; TYPE: PRT
; ORGANISM: *Homo sapiens*
US-11-169-041-145

Query Match 67.4%; Score 31; DB 11; Length 917;
Best Local Similarity 55.6%; Pred. No. 2.9e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

OY 1 ELQTTTHDI 9
|||
DB 214 ELQTTTHDI 222

RESULT 47

US-10-915-002-321
; Sequence 321, Application US/10915002
; Publication No. US20060078950A1
; GENERAL INFORMATION:
; APPLICANT: Proguiske-Fox, Ann
; APPLICANT: Hillman, Jeffrey D.
; TITLE OF INVENTION: IDENTIFICATION OF PORPHYROMONAS GINGIVALIS VIRULENCE POLYNUCLEOT
; TITLE OF INVENTION: USE IN DIAGNOSIS ANTIGENS FOR USE IN THE DIAGNOSIS, TREATMENT,
; TITLE OF INVENTION: PERIODONTAL DISEASES
; FILE REFERENCE: 02-042
; CURRENT APPLICATION NUMBER: US/10/915,002
; CURRENT FILING DATE: 2004-08-10
; NUMBER OF SEQ ID NOS: 354
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 321
; LENGTH: 1039
; TYPE: PRT
; ORGANISM: *Porphyromonas gingivalis*
US-10-915-002-321

Query Match 67.4%; Score 31; DB 9; Length 1039;
Best Local Similarity 33.3%; Pred. No. 3.4e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

OY 1 ELQTTTHDI 9
|||
DB 694 QIKTSVHDV 702

RESULT 48

US-10-915-002-322
; Sequence 322, Application US/10915002
; Publication No. US20060078950A1
; GENERAL INFORMATION:
; APPLICANT: Proguiske-Fox, Ann
; APPLICANT: Hillman, Jeffrey D.
; APPLICANT: Handfield, Martin
; TITLE OF INVENTION: IDENTIFICATION OF PORPHYROMONAS GINGIVALIS VIRULENCE POLYNUCLEOT
; TITLE OF INVENTION: USE IN DIAGNOSIS ANTIGENS FOR USE IN THE DIAGNOSIS, TREATMENT,
; FILE REFERENCE: 02-042
; CURRENT APPLICATION NUMBER: US/10/915,002
; CURRENT FILING DATE: 2004-08-10
; NUMBER OF SEQ ID NOS: 354

SOFTWARE: PatentIn version 3.1
SEQ ID NO 322
LENGTH: 1039
TYPE: PRT
ORGANISM: Porphyromonas gingivalis
US-10-915-002-322

Query Match 67.4%; Score 31; DB 9; Length 1039;
Best Local Similarity 33.3%; Pred. No. 3.4e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

QY 1 EL0TTIHD1 9
Db 694 QIKTSVHDV 702

RESULT 49
US-10-915-002-323
Sequence 323, Application US/10915002
Publication No. US20060078950A1
GENERAL INFORMATION:
APPLICANT: Proguiske-Fox, Ann
APPLICANT: Hillman, Jeffrey D.
APPLICANT: Handfield, Martin
TITLE OF INVENTION: IDENTIFICATION OF PORPHYROMONAS GINGIVALIS VIRULENCE POLYNUCLEOTI
TITLE OF INVENTION: USE IN DIAGNOSIS ANTIGENS FOR USE IN THE DIAGNOSIS, TREATMENT, A
TITLE OF INVENTION: PERIODONTAL DISEASES
FILE REFERENCE: 02-042
CURRENT APPLICATION NUMBER: US/10/915,002
CURRENT FILING DATE: 2004-08-10
NUMBER OF SEQ ID NOS: 354
SOFTWARE: PatentIn version 3.1
SEQ ID NO 323
LENGTH: 1039
TYPE: PRT
ORGANISM: Porphyromonas gingivalis
US-10-915-002-323

Query Match 67.4%; Score 31; DB 9; Length 1039;
Best Local Similarity 33.3%; Pred. No. 3.4e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

QY 1 EL0TTIHD1 9
Db 694 QIKTSVHDV 702

RESULT 50
US-11-096-568A-14692
Sequence 14692, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 14692
LENGTH: 1463
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(1463)
OTHER INFORMATION: Ceres Seq. ID no. 11049914
US-11-096-568A-14692

Query Match 67.4%; Score 31; DB 11; Length 1463;
Best Local Similarity 55.6%; Pred. No. 5e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 EL0TTIHD1 9
Db 858 KLOETLHDW 866

Search completed: May 5, 2006, 08:07:57
Job time : 10.5 secs

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 02:25:57 ; Search time 20.7 Seconds
(without alignments)
35.946 Million cell updates/sec

Title: US-08-170-344-6
Perfect score: 45
Sequence: 1 LQTTIHDI 9

Scoring table: BIOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%

Listing first 1000 summaries

Database : Issued Patents AA:*
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6: /cgn2_6/ptodata/1/1aa/backfile1.pep:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	* Query Match	Length	DB	ID	Description
1	45	100.0	9	1	US-08-787-547-102	Sequence 102, App
2	45	100.0	9	2	US-08-159-339A-248	Sequence 248, App
3	45	100.0	14	1	US-07-909-122-3	Sequence 4, Appli
4	45	100.0	30	1	US-08-363-586-4	Sequence 4, Appli
5	45	100.0	30	2	US-09-980-523A-4	Sequence 4, Appli
6	45	100.0	151	2	US-09-701-080C-18	Sequence 18, Appli
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26	33	73.3	297	2	US-09-097-231-8	Sequence 8, Appli
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265	28	62.2	102	4	PCT-US93-08742-12	Sequence 12, Appli	338	28	62.2	395	1	US-08-476-000-2	Sequence 2, Appli
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547	27	60.0	415	2	US-09-949-016-11269	Sequence 11269, A	620	27	60.0	931	2	US-10-037-417-118	Sequence 118, App
548	27	60.0	424	2	US-09-538-092-881	Sequence 881, App	621	27	60.0	931	2	US-10-037-417-119	Sequence 119, App
549	27	60.0	426	2	US-08-961-083-48	Sequence 48, Appl	622	27	60.0	931	2	US-10-037-417-120	Sequence 120, App
550	27	60.0	426	2	US-09-536-784-48	Sequence 48, Appl	623	27	60.0	967	2	US-09-139-802-201	Sequence 201, App
551	27	60.0	426	2	US-09-765-271-48	Sequence 48, Appl	624	27	60.0	967	2	US-09-659-786-201	Sequence 201, App
552	27	60.0	426	2	US-09-765-372A-48	Sequence 48, Appl	625	27	60.0	1031	2	US-09-306-595C-7	Sequence 7, Appl1
553	27	60.0	427	2	US-09-186-857-2	Sequence 2, Appl1	626	27	60.0	1031	2	US-09-925-388-7	Sequence 7, Appl1
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555	27	60.0	432	2	US-09-489-039A-8666	Sequence 8666, Ap	628	27	60.0	1123	2	US-10-037-417-71	Sequence 71, Appl
556	27	60.0	435	2	US-09-107-433-4256	Sequence 4256, Ap	629	27	60.0	1278	2	US-09-604-957-3	Sequence 3, Appl1
557	27	60.0	437	2	US-09-328-352-5193	Sequence 5193, Ap	630	27	60.0	1437	2	US-09-061-400-2	Sequence 2, Appl1
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564	27	60.0	475	2	US-09-328-352-4223	Sequence 4223, Ap	637	27	60.0	1781	2	US-09-995-749A-2	Sequence 2, Appl1
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568	27	60.0	485	2	US-10-130-419-1	Sequence 1, Appl1	641	27	60.0	2175	2	US-10-425-800-2	Sequence 2, Appl1
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570	27	60.0	507	2	US-09-134-001C-4892	Sequence 4892, Ap	643	27	60.0	2183	1	US-08-905-817-7	Sequence 7, Appl1
571	27	60.0	507	2	US-09-902-540-9980	Sequence 9980, Ap	644	27	60.0	2188	2	US-09-404-650-4	Sequence 4, Appl1
572	27	60.0	512	2	US-09-390-334-24	Sequence 24, Appl	645	27	60.0	2188	2	US-09-935-541-4	Sequence 4, Appl1
573	27	60.0	512	2	US-09-603-311-24	Sequence 30, Appl	646	27	60.0	2188	2	US-10-425-800-4	Sequence 4, Appl1
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578	27	60.0	560	2	US-09-487-370-6	Sequence 6, Appl1	651	27	60.0	7	2	US-09-479-130-4	Sequence 4, Appl1
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584	27	60.0	590	1	US-08-448-196A-9	Sequence 9, Appl1	657	27	60.0	42	2	US-09-480-993-20	Sequence 20, Appl
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590	27	60.0	623	2	US-09-608-533A-6	Sequence 6, Appl1	663	27	60.0	56	2	US-09-513-999C-6934	Sequence 6934, Ap
591	27	60.0	625	2	US-09-661-322A-48	Sequence 48, Appl	664	27	60.0	57	2	US-09-079-030-107	Sequence 107, App
592	27	60.0	632	2	US-09-661-322A-2	Sequence 2, Appl1	665	27	60.0	70	2	US-09-489-039A-7808	Sequence 7808, Ap
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597	27	60.0	634	2	US-09-186-002-12	Sequence 12, Appl	670	27	60.0	74	2	US-09-206-551-35	Sequence 35, Appl1
598	27	60.0	635	2	US-09-041-991A-4	Sequence 4, Appl1	671	27	60.0	74	2	US-09-134-000C-6520	Sequence 6520, Ap
599	27	60.0	635	2	US-09-608-533A-4	Sequence 4, Appl1	672	27	60.0	76	2	US-09-134-001C-4377	Sequence 4377, Ap
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607	27	60.0	713	1	US-08-987-466-3	Sequence 3, Appl1	680	27	60.0	86	1	US-08-461-859-20	Sequence 20, Appl
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686	26	57.8	107	1	US-08-446-363-5	Sequence 5, Appl1	759	26	57.8	260	2	US-09-443-041A-10	Sequence 10, Appl
687	26	57.8	114	2	US-09-489-847-321	Sequence 321, App	760	26	57.8	261	2	US-09-443-041A-26	Sequence 26, Appl
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693	26	57.8	142	2	US-08-906-769-155	Sequence 155, App	766	26	57.8	275	2	US-08-836-134-21	Sequence 12, Appl
694	26	57.8	142	2	US-08-906-616-155	Sequence 155, App	767	26	57.8	275	2	US-09-493-784-21	Sequence 12, Appl
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702	26	57.8	154	2	US-09-345-236B-144	Sequence 144, App	775	26	57.8	294	2	US-09-134-000C-4825	Sequence 30737, A
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706	26	57.8	159	2	US-09-640-211A-1023	Sequence 4756, Ap	779	26	57.8	312	2	US-09-270-767-32060	Sequence 32060, A
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712	26	57.8	164	2	US-09-270-767-51721	Sequence 1075, Ap	785	26	57.8	320	2	US-09-248-796A-17608	Sequence 17608, A
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717	26	57.8	180	2	US-09-543-681A-5017	Sequence 211, App	790	26	57.8	333	2	US-08-961-083-54	Sequence 54, Appl
718	26	57.8	180	2	US-09-079-030-211	Sequence 725, App	791	26	57.8	333	2	US-09-338-420-4	Sequence 4, Appl1
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722	26	57.8	190	2	US-09-248-796A-26836	Sequence 26836, A	795	26	57.8	333	2	US-09-765-272A-54	Sequence 14, Appl
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725	26	57.8	202	2	US-10-104-047-2736	Sequence 2736, Ap	798	26	57.8	336	2	US-08-273-247-2	Sequence 12, Appl
726	26	57.8	210	2	US-09-079-030-115	Sequence 115, App	799	26	57.8	336	2	US-09-878-766A-12	Sequence 14, Appl
727	26	57.8	210	2	US-09-583-110-3821	Sequence 2821, Ap	800	26	57.8	336	2	US-09-878-766A-14	Sequence 16, Appl
728	26	57.8	214	2	US-09-902-540-16021	Sequence 16021, A	801	26	57.8	336	2	US-09-878-766A-16	Sequence 18, Appl
729	26	57.8	216	2	US-09-345-236B-115	Sequence 115, App	802	26	57.8	336	2	US-09-878-766A-18	Sequence 20, Appl
730	26	57.8	217	2	US-09-107-433-3537	Sequence 3537, App	803	26	57.8	336	2	US-09-878-766A-20	Sequence 4, Appl1
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734	26	57.8	224	2	US-09-465-901-30	Sequence 30, Appl	807	26	57.8	336	2	US-10-134-297-10	Sequence 12, Appl
735	26	57.8	231	2	US-08-311-731A-74	Sequence 74, Appl	808	26	57.8	336	2	US-10-134-297-12	Sequence 14, Appl
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738	26	57.8	242	2	US-09-004-731-36	Sequence 36, Appl	811	26	57.8	336	2	US-09-878-781-8	Sequence 8, Appl1
739	26	57.8	242	2	US-08-749-699-36	Sequence 36, Appl	812	26	57.8	336	2	US-09-878-781-10	Sequence 10, Appl
740	26	57.8	242	2	US-09-004-729-36	Sequence 22943, A	813	26	57.8	336	2	US-09-878-781-12	Sequence 12, Appl
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745	26	57.8	244	2	US-09-583-110-4115	Sequence 4115, Ap	818	26	57.8	336	2	US-10-650-369-18	Sequence 18, Appl
746	26	57.8	249	2	US-09-540-236-3796	Sequence 3796, Ap	819	26	57.8	336	2	US-10-650-369-20	Sequence 20, Appl
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749	26	57.8	256	2	US-09-107-433-3533	Sequence 3533, App	822	26	57.8	340	2	US-09-252-991A-21612	Sequence 21612, A
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755	26	57.8	259	2	US-08-749-699-85	Sequence 190, App	828	26	57.8	349	2	US-09-543-681A-8004	Sequence 8004, Ap
756	26	57.8	259	2	US-09-012-692-190	Sequence 190, App	829	26	57.8	352	2	US-09-198-452A-582	Sequence 582, App
757	26	57.8	259	2	US-08-906-613-190	Sequence 190, App	830	26	57.8				

831	26	57.8	353	1	US-08-118-270-45	Sequence 45, Appl	904	26	57.8	454	2	US-09-905-589-11	Sequence 11, Appl
832	26	57.8	353	1	US-08-176-620A-14	Sequence 14, Appl	905	26	57.8	454	2	US-10-108-171A-11	Sequence 11, Appl
833	26	57.8	353	1	US-08-461-985-14	Sequence 14, Appl	906	26	57.8	456	2	US-09-543-681A-5230	Sequence 5230, Ap
834	26	57.8	353	2	US-08-932-787B-19	Sequence 19, Appl	907	26	57.8	458	2	US-09-134-001C-4663	Sequence 4663, Ap
835	26	57.8	353	2	US-08-932-012C-19	Sequence 19, Appl	908	26	57.8	458	2	US-09-902-540-16600	Sequence 16600, A
836	26	57.8	353	2	US-08-888-818C-19	Sequence 19, Appl	909	26	57.8	463	2	US-09-163-44A-2	Sequence 2, Appl1
837	26	57.8	353	2	US-09-614-912-204	Sequence 204, App	910	26	57.8	465	2	US-09-902-540-13166	Sequence 13166, A
838	26	57.8	353	2	US-09-487-558B-116	Sequence 116, App	911	26	57.8	476	2	US-10-214-269-2	Sequence 2, Appl1
839	26	57.8	353	4	PCT-US93-08528-45	Sequence 45, Appl	912	26	57.8	479	2	US-09-252-991A-23370	Sequence 23370, A
840	26	57.8	356	1	US-08-681-151-1	Sequence 1, Appl1	913	26	57.8	482	2	US-09-438-185A-356	Sequence 356, App
841	26	57.8	358	1	US-09-438-185A-546	Sequence 546, App	914	26	57.8	487	2	US-09-134-000C-6001	Sequence 6001, Ap
842	26	57.8	359	2	US-09-108-020-45	Sequence 45, Appl	915	26	57.8	492	2	US-09-999-833A-7	Sequence 7, Appl1
843	26	57.8	359	2	US-09-583-110-4722	Sequence 4722, Ap	916	26	57.8	492	2	US-10-020-448A-7	Sequence 7, Appl1
844	26	57.8	359	2	US-09-685-296-45	Sequence 45, Appl	917	26	57.8	493	2	US-09-489-039A-12903	Sequence 12903, A
845	26	57.8	361	2	US-09-540-236-3112	Sequence 3112, Ap	918	26	57.8	496	1	US-08-838-543-2	Sequence 2, Appl1
846	26	57.8	362	2	US-09-596-824-6	Sequence 6, Appl1	919	26	57.8	497	1	US-08-740-223A-14	Sequence 14, Appl1
847	26	57.8	362	2	US-09-885-329-6	Sequence 6, Appl1	920	26	57.8	497	2	US-09-709-188-14	Sequence 14, Appl
848	26	57.8	365	2	US-09-748-264A-2	Sequence 2, Appl1	921	26	57.8	497	2	US-10-225-060-14	Sequence 14, Appl
849	26	57.8	366	2	US-09-596-824-4	Sequence 4, Appl1	922	26	57.8	507	2	US-10-214-269-14	Sequence 14, Appl
850	26	57.8	366	2	US-09-885-329-4	Sequence 4, Appl1	923	26	57.8	508	2	US-09-270-767-46718	Sequence 46718, A
851	26	57.8	366	2	US-09-489-039A-10346	Sequence 10346, A	924	26	57.8	518	2	US-09-252-991A-25967	Sequence 25967, A
852	26	57.8	366	2	US-09-270-767-43543	Sequence 43543, A	925	26	57.8	544	2	US-09-489-039A-14296	Sequence 14296, A
853	26	57.8	367	2	US-09-248-796A-18976	Sequence 18976, A	926	26	57.8	549	2	US-10-104-047-3479	Sequence 3479, Ap
854	26	57.8	368	2	US-09-248-796A-17319	Sequence 17319, A	927	26	57.8	557	2	US-09-808-387-10	Sequence 10, Appl
855	26	57.8	369	2	US-09-107-433-2790	Sequence 2790, Ap	928	26	57.8	558	2	US-09-710-279-1760	Sequence 1760, Ap
856	26	57.8	381	2	US-09-902-540-11854	Sequence 11854, A	929	26	57.8	564	2	US-09-107-532A-6235	Sequence 6235, Ap
857	26	57.8	383	2	US-09-134-001C-3701	Sequence 3701, Ap	930	26	57.8	564	2	US-09-107-532A-6237	Sequence 6237, Ap
858	26	57.8	383	2	US-09-638-937-7	Sequence 7, Appl1	931	26	57.8	567	1	US-08-841-483-2	Sequence 2, Appl1
859	26	57.8	383	2	US-09-710-279-3110	Sequence 3110, Ap	932	26	57.8	567	2	US-09-382-911-2	Sequence 8, Appl1
860	26	57.8	386	2	US-09-338-420-2	Sequence 2, Appl1	933	26	57.8	572	2	US-09-648-004-8	Sequence 8, Appl1
861	26	57.8	386	2	US-09-248-796A-16103	Sequence 16103, A	934	26	57.8	572	2	US-10-272-419-8	Sequence 8, Appl1
862	26	57.8	387	1	US-08-837-593-4	Sequence 4, Appl1	935	26	57.8	581	2	US-09-107-532A-6835	Sequence 6835, Ap
863	26	57.8	387	2	US-09-032-215-8	Sequence 8, Appl1	936	26	57.8	584	2	US-08-753-750B-4	Sequence 4, Appl1
864	26	57.8	387	2	US-09-032-215-13	Sequence 13, Appl	937	26	57.8	593	2	US-09-252-991A-55274	Sequence 25274, A
865	26	57.8	388	2	US-09-502-540-14227	Sequence 14227, A	938	26	57.8	594	2	US-09-380-287A-8	Sequence 8, Appl1
866	26	57.8	391	2	US-09-902-540-12443	Sequence 12443, A	939	26	57.8	594	2	US-09-380-287A-10	Sequence 10, Appl
867	26	57.8	393	1	US-08-837-593-3	Sequence 3, Appl1	940	26	57.8	595	2	US-09-248-796A-14562	Sequence 14562, A
868	26	57.8	395	2	US-09-949-016-8564	Sequence 8564, Ap	941	26	57.8	599	2	US-09-252-991A-20656	Sequence 20656, A
869	26	57.8	400	2	US-09-004-731-30	Sequence 30, Appl	942	26	57.8	601	1	US-08-333-358-14	Sequence 14, Appl
870	26	57.8	400	2	US-09-004-731-33	Sequence 33, Appl	943	26	57.8	601	1	US-08-463-694-14	Sequence 14, Appl
871	26	57.8	400	2	US-08-749-699-30	Sequence 30, Appl	944	26	57.8	601	1	US-08-694-501-14	Sequence 14, Appl
872	26	57.8	400	2	US-08-749-699-33	Sequence 33, Appl	945	26	57.8	608	2	US-09-134-000C-5687	Sequence 5687, Ap
873	26	57.8	400	2	US-09-004-729-30	Sequence 30, Appl	946	26	57.8	612	2	US-09-569-611C-50	Sequence 50, Appl
874	26	57.8	400	2	US-09-004-729-33	Sequence 33, Appl	947	26	57.8	621	2	US-09-720-317A-16	Sequence 16, Appl
875	26	57.8	400	2	US-09-270-767-46136	Sequence 46136, A	948	26	57.8	635	2	US-09-248-796A-23137	Sequence 23137, A
876	26	57.8	401	2	US-09-522-991A-27198	Sequence 27198, A	949	26	57.8	636	1	US-08-248-796A-14866	Sequence 14866, A
877	26	57.8	402	2	US-09-025-578-2	Sequence 2, Appl1	950	26	57.8	646	1	US-08-441-139-14	Sequence 14, Appl
878	26	57.8	402	2	US-09-330-235-2	Sequence 2, Appl1	951	26	57.8	646	2	US-09-919-039-11	Sequence 11, Appl
879	26	57.8	402	2	US-10-104-339-2	Sequence 2, Appl1	952	26	57.8	656	2	US-09-327-984A-36	Sequence 36, Appl
880	26	57.8	403	2	US-09-248-796A-19438	Sequence 19438, A	953	26	57.8	658	2	US-08-953-040-9	Sequence 9, Appl1
881	26	57.8	406	1	US-09-134-001C-3202	Sequence 3202, Ap	954	26	57.8	676	2	US-09-107-532A-5627	Sequence 5627, Ap
882	26	57.8	408	1	US-08-742-440A-6	Sequence 6, Appl1	955	26	57.8	678	1	US-08-288-408-5	Sequence 5, Appl1
883	26	57.8	412	2	US-09-489-039A-12933	Sequence 12933, A	956	26	57.8	678	1	US-08-655-782-5	Sequence 5, Appl1
884	26	57.8	415	2	US-09-710-279-1406	Sequence 1406, Ap	957	26	57.8	701	2	US-08-923-511-2	Sequence 2, Appl1
885	26	57.8	417	2	US-09-902-540-10785	Sequence 10785, A	958	26	57.8	701	2	US-09-415-874A-2	Sequence 2, Appl1
886	26	57.8	418	2	US-09-743-742B-5	Sequence 5, Appl1	959	26	57.8	713	2	US-09-270-767-45101	Sequence 45101, A
887	26	57.8	418	2	US-09-826-509-535	Sequence 535, Appl	960	26	57.8	713	2	US-09-902-540-16350	Sequence 16350, A
888	26	57.8	421	2	US-09-540-236-2339	Sequence 2339, Ap	961	26	57.8	715	2	US-09-538-092-970	Sequence 970, App
889	26	57.8	425	2	US-09-489-039A-7338	Sequence 7338, Ap	962	26	57.8	718	1	US-08-974-565C-7	Sequence 7, Appl1
890	26	57.8	425	2	US-09-540-236-2974	Sequence 2974, Ap	963	26	57.8	718	2	US-09-255-748-7	Sequence 7, Appl1
891	26	57.8	427	2	US-09-689-065B-3	Sequence 3, Appl1	964	26	57.8	727	2	US-09-487-558B-296	Sequence 296, App
892	26	57.8	432	2	US-09-583-110-5039	Sequence 5039, Ap	965	26	57.8	733	2	US-09-192-983-6	Sequence 6, Appl1
893	26	57.8	433	2	US-09-107-433-3456	Sequence 3456, Ap	966	26	57.8	737	2	US-09-955-732A-13	Sequence 13, Appl
894	26	57.8	439	2	US-09-328-352-4368	Sequence 4368, Ap	967	26	57.8	738	2	US-09-248-796A-16666	Sequence 16666, A
895	26	57.8	448	2	US-09-878-766A-22	Sequence 22, Appl	968	26	57.8	739	2	US-09-902-540-9801	Sequence 9801, Ap
896	26	57.8	448	2	US-10-650-369-22	Sequence 22, Appl	969	26	57.8	748	2	US-08-904-871-6	Sequence 6, Appl1
897	26	57.8	454	2	US-09-240-639-11	Sequence 11, Appl	970	26	57.8	748	2	US-08-904-871-13	Sequence 13, Appl
898	26	57.8	454	2	US-09-908-510A-11	Sequence 11, Appl	971	26	57.8	752	1	US-08-309-512-8	Sequence 8, Appl1
899	26	57.8	454	2	US-09-905-744B-11	Sequence 11, Appl	972	26	57.8	752	4	PCT-US92-08756A-8	Sequence 8, Appl1
900	26	57.8	454	2	US-10-107-660-11	Sequence 11, Appl	973	26	57.8	763	1	US-09-538-092-684	Sequence 684, App
901	26	57.8	454	2	US-10-107-576-11	Sequence 11, Appl	974	26	57.8	768	2	US-09-248-796A-20025	Sequence 20025, A
902	26	57.8	454	2	US-09-905-733B-11	Sequence 11, Appl	975	26	57.8	795	2	US-09-328-352-6143	Sequence 6143, Ap
903	26	57.8	454	2	US-09-905-743B-11	Sequence 11, Appl	976	26	57.8	801	2	US-09-913-301-7	Sequence 7, Appl1

977 26 57.8 807 2 US-09-177-650-3
978 26 57.8 822 1 US-08-222-617A-7
979 26 57.8 825 2 US-09-328-352-5974
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982 26 57.8 872 2 US-09-949-016-8039
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985 26 57.8 872 2 US-09-949-016-8042
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987 26 57.8 876 2 US-08-996-621-2
988 26 57.8 883 1 US-08-953-492-2
989 26 57.8 883 2 US-09-583-110-2900
990 26 57.8 883 2 US-09-248-796A-14418
991 26 57.8 888 2 US-09-134-001C-3032
992 26 57.8 888 2 US-09-107-433-2964
993 26 57.8 890 2 US-09-513-783A-174
994 26 57.8 890 2 US-10-100-957A-174
995 26 57.8 894 2 US-09-949-016-10605
996 26 57.8 898 1 US-08-808-982-5
997 26 57.8 898 2 US-09-306-902A-5
998 26 57.8 899 2 US-09-540-236-3664
999 26 57.8 906 2 US-09-134-001C-3218
1000 26 57.8 922 2 US-09-902-540-12187

ALIGNMENTS

Sequence 3, Appl
Sequence 7, Appl
Sequence 5974, Ap
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Sequence 8040, Ap
Sequence 8041, Ap
Sequence 8042, Ap
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Sequence 2, Appl
Sequence 2900, Ap
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Sequence 2964, Ap
Sequence 174, App
Sequence 174, App
Sequence 10605, A
Sequence 5, Appl
Sequence 5, Appl
Sequence 3664, Ap
Sequence 3218, Ap
Sequence 12187, A

RESULT 1
US-08-787-547-102
Sequence 102, Application US/08787547
Patent No. 5783567
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Curley, Joanne M.
APPLICANT: Langer, Robert S.
TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY
TITLE OF INVENTION: OF NUCLEIC ACID
NUMBER OF SEQUENCES: 107
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/787,547
FILING DATE: 22-JAN-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
ATTORNEY/AGENT INFORMATION:
FILING DATE:
NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08191/003001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 102:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide

US-08-787-547-102

Query Match 100.0%; Score 45; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 1 LOTTTHDII 9
Db 1 LOTTTHDII 9

RESULT 2
US-08-159-339A-248
Sequence 248, Application US/08159339A
Patent No. 6037135

GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esben
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lawyer
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:

INFORMATION FOR SEQ ID NO: 248:
SEQUENCE CHARACTERISTICS:

LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-248

Query Match 100.0%; Score 45; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 1 LOTTTHDII 9
Db 1 LOTTTHDII 9

RESULT 3

US-07-909-122-3
; Sequence 3, Application US/07909122
; Patent No. 5415995
; GENERAL INFORMATION:
; APPLICANT: SCHOOLNIK, GARY K.
; APPLICANT: PALERSKY, JOEL M.
; TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
; TITLE OF INVENTION: VIRUS
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MORRISON & FOERSTER
; STREET: 755 Page Mill Road
; CITY: Palo Alto
; STATE: California
; COUNTRY: USA
; ZIP: 94304-1018
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/07/909,122
; FILING DATE: 19920706
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: BENZ, WILLIAM H.
; REGISTRATION NUMBER: 25,952
; REFERENCE/DOCKET NUMBER: 28600-20105.01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 813-5600
; TELEFAX: (415) 494-0792
; TELEX: 706141
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 14 amino acids
; TYPE: AMINO ACID
; STRANDEDNESS: single
; TOPOLOGY: linear
; US-07-909-122-3

Query Match 100.0%; Score 45; DB 1; Length 14;
Best Local Similarity 100.0%; Pred. No. 0.036;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
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| | | | |
Db 3 LQTTIHDI 11

RESULT 4
US-08-363-586-4
; Sequence 4, Application US/08363586
; Patent No. 5629161
; GENERAL INFORMATION:
; APPLICANT: Mueller, Martin
; APPLICANT: Giesmann, Lutz
; TITLE OF INVENTION: Use of HPV-16 B6 and E7-Gene Derived
; TITLE OF INVENTION: Peptides for the Diagnostic Purpose
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
; ADDRESSEE: Dunner
; STREET: 1300 I Street, N.W.
; CITY: Washington
; STATE: D.C.
; COUNTRY: USA
; ZIP: 20005-3315
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25

US-08-363-586-4
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/363,586
; FILING DATE: 23-DEC-1994
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/909,296
; FILING DATE: 09-JUL-1992
; APPLICATION NUMBER: EP 91111720.8
; FILING DATE: 13-JUL-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Wadler, Linda A.
; REGISTRATION NUMBER: 33,218
; REFERENCE/DOCKET NUMBER: 02481-1195-00000
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 202-408-4400
; TELEFAX: 202-408-4400
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 30 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; US-08-363-586-4

Query Match 100.0%; Score 45; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.08;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
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Db 19 LQTTIHDI 27

RESULT 5
US-09-980-523A-4
; Sequence 4, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: PERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE B6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: WO/1 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO: 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
; US-09-980-523A-4

Query Match 100.0%; Score 45; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.08;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
| | | | |
| | | | |
Db 12 LQTTIHDI 20

RESULT 6
US-09-701-080C-18
; Sequence 18, Application US/09701080C


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; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; PRIOR FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
;
US-09-701-080C-18
;
Query Match          100.0%; Score 45; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.43;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 LQTTIHDI 9
        |||||
        19 LQTTIHDI 27

RESULT 7
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCES
; APPLICANT: FERRIS, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: WO/01 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
;
US-09-980-523A-2
;
Query Match          100.0%; Score 45; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.46;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 LQTTIHDI 9
        |||||
        26 LQTTIHDI 34

RESULT 8
US-08-316-239B-3
; Sequence 3, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
;

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; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
;
US-08-316-239B-3
;
Query Match          100.0%; Score 45; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 LQTTIHDI 9
        |||||
        26 LQTTIHDI 34

RESULT 9
US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
;

```


FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtland, Ajay A.
REGISTRATION NUMBER: 35, 205
REFERENCE/DOCKET NUMBER: UNNE-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-4

Query Match 100.0%; Score 45; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.47;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDTI 9
|||
Db 26 LQTTIHDTI 34

RESULT 10
US-08-860-165-14
Sequence 14, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860.165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14

Query Match 100.0%; Score 45; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDTI 9
|||
Db 95 LQTTIHDTI 103

RESULT 11
US-09-359-382-14
Sequence 14, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian

TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359.382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860.165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-14

Query Match 100.0%; Score 45; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDTI 9
|||
Db 95 LQTTIHDTI 103

RESULT 12
US-08-117-083-10
Sequence 10, Application US/08117083
Patent No. 5719054
GENERAL INFORMATION:
APPLICANT: Bournselli, Michael E.
APPLICANT: Inglis, Stephen C.
APPLICANT: Munro, Alan J.
TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSER: Walter H. Dregger
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117.083
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dregger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-58783
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 182 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..182
OTHER INFORMATION: /note= "Xaa refers to stop codon in

OTHER INFORMATION: the open reading frame."

US-08-117-083-10
Query Match 100.0%; Score 45; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.53;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 27 LQTTIHDI 35

RESULT 13
US-09-462-993-1
Sequence 1, Application US/09462993
Patent No. 6884786
GENERAL INFORMATION:
APPLICANT: KIENY, Marie-Paule
APPLICANT: BALLOU, Jean-Marc
APPLICANT: BIZOUARNE, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
POLYPEPTIDE WITH MODIFIED CELL LOCATION
FILE REFERENCE: 017753-122
CURRENT APPLICATION NUMBER: US/09/462,993
CURRENT FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: PCT/FR98/01576
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
human papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-09-462-993-1

Query Match 100.0%; Score 45; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 54 LQTTIHDI 62

RESULT 14
US-08-860-165-10
Sequence 10, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
PRIOR APPLICATION NUMBER: PCT/AU95/00868
PRIOR FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Artificial Sequence

FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10

Query Match 100.0%; Score 45; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.79;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 26 LQTTIHDI 34

RESULT 15
US-09-359-382-10
Sequence 10, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
PRIOR APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
PRIOR APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 45; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.79;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 26 LQTTIHDI 34

RESULT 16
US-09-367-309A-1
Sequence 1, Application US/09367309A
Patent No. 6428807
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 45; DB 2; Length 266;

Best Local Similarity 100.0%; Pred. No. 0.79; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;
OY 1 LQTTIHDI 9
|||
Db 26 LQTTIHDI 34

RESULT 17
US-09-485-885-4
; Sequence 4, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 45; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.81; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;
OY 1 LQTTIHDI 9
|||
Db 132 LQTTIHDI 140

RESULT 18
US-09-485-885-10
; Sequence 10, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 45; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.87; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

OY 1 LQTTIHDI 9
|||
Db 151 LQTTIHDI 159

RESULT 19
US-09-485-885-6
; Sequence 6, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 45; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 1.1; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;
OY 1 LQTTIHDI 9
|||
Db 132 LQTTIHDI 140

RESULT 20
US-09-485-885-14
; Sequence 14, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 45; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 1.2; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;
OY 1 LQTTIHDI 9
|||
Db 151 LQTTIHDI 159

RESULT 21
US-08-934-915-159
; Sequence 159, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILNER, JOAKIM
; APPLICANT: DILNER, LENA
; APPLICANT: CHENG, HMEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
; NUMBER OF SEQUENCES: 193
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft Word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. Fouch
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946.6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:
; INFORMATION FOR SEQ ID NO: 159:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; US-08-934-915-159

Query Match 88.9%; Score 40; DB 1; Length 20;
Best Local Similarity 88.9%; Pred. No. 0.44;
Matches 8; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 10 LQTTIHNI 18

RESULT 22
US-08-913-159-14
; Sequence 14, Application US/08913159
; Patent No. 6300109
; GENERAL INFORMATION:
; APPLICANT:
; TITLE OF INVENTION: Plasmid-derived type II
; TITLE OF INVENTION: restriction-modification systems from Lactococcus lactis
; NUMBER OF SEQUENCES: 14
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/913,159
; FILING DATE:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: DK 0179/95
; FILING DATE: 17-FEB-1995
; INFORMATION FOR SEQ ID NO: 14:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 317 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; US-08-913-159-14

Query Match 88.9%; Score 40; DB 2; Length 317;
Best Local Similarity 77.8%; Pred. No. 8;
Matches 7; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 182 LKTTIHDI 190

RESULT 23
US-09-270-767-33046
; Sequence 33046, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 33046
; LENGTH: 166
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; US-09-270-767-33046

Query Match 73.3%; Score 33; DB 2; Length 166;
Best Local Similarity 62.5%; Pred. No. 81;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 8
Db 85 LRTTIHDL 92

RESULT 24
US-09-270-767-48263
; Sequence 48263, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 48263
; LENGTH: 166
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; US-09-270-767-48263

Query Match 73.3%; Score 33; DB 2; Length 166;
Best Local Similarity 62.5%; Pred. No. 81;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 8

Db :||:|:
85 LRTLHDL 92

RESULT 25
US-08-706-281A-8
Sequence 8, Application US/08706281A
Patent No. 6100048
GENERAL INFORMATION:
APPLICANT: Cone, Roger D
APPLICANT: Fan, Wei
APPLICANT: Boston, Bruce A
APPLICANT: Kesterton, Robert A
APPLICANT: Lu, Dongxi
APPLICANT: Chen, Wendiao
TITLE OF INVENTION: Methods and Reagents for Discovering and
TITLE OF INVENTION: Using Mammalian Melanocortin Receptor Agonists and Antagonists
TITLE OF INVENTION: To Modulate Feeding Behavior in Animals
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSER: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive
CITY: Chicago
STATE: IL
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/706/281A
FILING DATE: 04-SEP-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: No. 6100048nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 96,886
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-0002
TELEX:
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 297 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-706-281A-8

Query Match 73.3%; Score 33; DB 2; Length 297;
Best Local Similarity 75.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 QTTIHDI 9
: || || ||
Db 99 ETTAHDI 106

RESULT 26
US-09-097-231-8
Sequence 8, Application US/09097231
Patent No. 6278038
GENERAL INFORMATION:
APPLICANT: Cone, Roger D
APPLICANT: Low, Malcolm J
APPLICANT: Chen, Wendiao
TITLE OF INVENTION: Mammalian Melanocortin Receptor and Uses
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSER: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive

CITY: Chicago
STATE: Illinois
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/097,231
FILING DATE: 12-Jun-1998
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: No. 6278038nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 96,886-C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-0002
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 297 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-097-231-8

Query Match 73.3%; Score 33; DB 2; Length 297;
Best Local Similarity 75.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 2 QTTIHDI 9
: || || ||
Db 99 ETTAHDI 106

RESULT 27
US-09-353-099-8
Sequence 8, Application US/09353099
Patent No. 6476187
GENERAL INFORMATION:
APPLICANT: Cone, Roger D
APPLICANT: Fan, Wei
APPLICANT: Boston, Bruce A
APPLICANT: Kesterton, Robert A
APPLICANT: Lu, Dongxi
APPLICANT: Chen, Wendiao
TITLE OF INVENTION: Methods and Reagents for Discovering and
TITLE OF INVENTION: Using Mammalian Melanocortin Receptor Agonists and Antagoni
TITLE OF INVENTION: To Modulate Feeding Behavior in Animals
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
ADDRESSER: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive
CITY: Chicago
STATE: IL
COUNTRY: USA
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/353,099
FILING DATE: 14-Sep-1999
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/706,281
FILING DATE: 04-SEP-1996

```

ATTORNEY/AGENT INFORMATION:
NAME: No. 6476187nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 96,886
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-0002
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 297 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-353-099-8

Query Match 73.3%; Score 33; DB 2; Length 297;
Best Local Similarity 100.0%; Pred. No. 1.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 QTTIID 9
DB 99 ETTAHDII 106

RESULT 28
US-09-354-123-6
Sequence 6, Application US/09354123
Patent No. 6489537
GENERAL INFORMATION:
APPLICANT: Rea, Philip A.
APPLICANT: Valamianuk, Olena K.
APPLICANT: Mari, Stephane
APPLICANT: Lu, Yu-Ping
APPLICANT: Schroeder, Julian I.
APPLICANT: Kim, Eugene J.
APPLICANT: Clemens, Stephan
TITLE OF INVENTION: NOVEL PHYTOCHELATIN SYNTHASES AND USES THEREFOR
FILE REFERENCE: 9596-10201/209596.0289
CURRENT APPLICATION NUMBER: US/09/354,123
CURRENT FILING DATE: 1999-07-15
EARLIER APPLICATION NUMBER: 09/315,449
EARLIER FILING DATE: 1999-05-20
EARLIER APPLICATION NUMBER: 60/095,624
EARLIER FILING DATE: 1998-08-07
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 500
TYPE: PRT
ORGANISM: Trifolium aestivum
US-09-354-123-6

Query Match 73.3%; Score 33; DB 2; Length 500;
Best Local Similarity 100.0%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 QTTIID 7
DB 125 QTTIHD 130

RESULT 29
US-10-214-269-19
Sequence 19, Application US/10214269
Patent No. 6844485
GENERAL INFORMATION:
APPLICANT: Butler, Karlene H.
APPLICANT: Famodu, Omolayo O.
APPLICANT: Harvell, Leslie T.
APPLICANT: Orozco, Jr., Emil M.
APPLICANT: Rasco-Gaunt, Sonliza

ATTORNEY/AGENT INFORMATION:
NAME: No. 6476187nan, Kevin E
REGISTRATION NUMBER: 35,303
REFERENCE/DOCKET NUMBER: 96,886
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312-913-0001
TELEFAX: 312-913-0002
TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 297 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 8:
US-09-353-099-8

Query Match 73.3%; Score 33; DB 2; Length 500;
Best Local Similarity 100.0%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 2 QTTIID 7
DB 125 QTTIHD 130

RESULT 30
US-09-270-767-32894
Sequence 32894, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 32894
LENGTH: 168
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-32894

Query Match 71.1%; Score 32; DB 2; Length 168;
Best Local Similarity 50.0%; Pred. No. 1.3e+02;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 2 QTTIID 9
DB 134 QATLHDVV 141

RESULT 31
US-09-489-039A-14156
Sequence 14156, Application US/09489039A
Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
FILE REFERENCE: 2709,2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 14156
LENGTH: 467
TYPE: PRT
ORGANISM: Klebsiella pneumoniae
US-09-489-039A-14156

Query Match 71.1%; Score 32; DB 2; Length 467;
Best Local Similarity 83.3%; Pred. No. 3.7e+02;

```

Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 2 QTTIHD 7
|||:|
Db 242 QTTVHD 247

RESULT 32
US-09-769-787-37
; Sequence 37, Application US/09769787
; Patent No. 6936252
; GENERAL INFORMATION:
; APPLICANT: Microbial Technics Limited
; APPLICANT: Gilbert, Christophe FG
; APPLICANT: Hansbro, Philip M
; TITLE OF INVENTION: Proteins
; FILE REFERENCE: PMC/P21129WO
; CURRENT APPLICATION NUMBER: US/09/769,787
; CURRENT FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: GB 9816337.1
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: US 60/125164
; PRIOR FILING DATE: 1999-03-19
; NUMBER OF SEQ ID NOS: 388
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 37
; LENGTH: 513
; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-769-787-37

Query Match 71.1%; Score 32; DB 2; Length 513;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTIHDI 8
|||||
Db 495 TTIHDI 500

RESULT 33
US-09-769-787-193
; Sequence 193, Application US/09769787
; Patent No. 6936252
; GENERAL INFORMATION:
; APPLICANT: Microbial Technics Limited
; APPLICANT: Gilbert, Christophe FG
; APPLICANT: Hansbro, Philip M
; TITLE OF INVENTION: Proteins
; FILE REFERENCE: PMC/P21129WO
; CURRENT APPLICATION NUMBER: US/09/769,787
; CURRENT FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: GB 9816337.1
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: US 60/125164
; PRIOR FILING DATE: 1999-03-19
; NUMBER OF SEQ ID NOS: 388
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 193
; LENGTH: 513
; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-09-769-787-193

Query Match 71.1%; Score 32; DB 2; Length 513;
Best Local Similarity 100.0%; Pred. No. 4.1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTIHDI 8
|||||
Db 495 TTIHDI 500

RESULT 34
US-09-252-991A-30492
; Sequence 30492, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 30492
; LENGTH: 532
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-30492

Query Match 71.1%; Score 32; DB 2; Length 532;
Best Local Similarity 55.6%; Pred. No. 4.2e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 LQTTIHDII 9
|||:|:|
Db 163 LQTTFHDIIV 171

RESULT 35
US-09-543-681A-6825
; Sequence 6825, Application US/09543681A
; Patent No. 6605709
; GENERAL INFORMATION:
; APPLICANT: GARY BRETON
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PROTEUS MIRABILIS
; TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 2709.1002-001
; CURRENT APPLICATION NUMBER: US/09/543,681A
; CURRENT FILING DATE: 2000-04-05
; PRIOR APPLICATION NUMBER: US 60/128,706
; PRIOR FILING DATE: 1999-04-09
; NUMBER OF SEQ ID NOS: 8344
; SEQ ID NO 6825
; LENGTH: 572
; TYPE: PRT
; ORGANISM: Proteus mirabilis
US-09-543-681A-6825

Query Match 71.1%; Score 32; DB 2; Length 572;
Best Local Similarity 71.4%; Pred. No. 4.6e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 QTTIHDII 8
|||:|
Db 406 ETTIHDII 412

RESULT 36
US-09-248-796A-19137
; Sequence 19137, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstein et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13

PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 19137
LENGTH: 781
TYPE: PRT
ORGANISM: Candida albicans
US-09-248-796A-19137

Query Match 71.1%; Score 32; DB 2; Length 781;
Best Local Similarity 62.5%; Pred. No. 6.3e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 QTHDII 9
DB 75 QTHLHDLI 82

RESULT 37
US-08-197-484-72
Sequence 72, Application US/08197484
Patent No. 6419931
GENERAL INFORMATION:
APPLICANT: VITIELLO, Maria A.
APPLICANT: CHESTNUT, Robert W.
APPLICANT: SETTE, Alessandro D.
APPLICANT: CELIS, Esben
APPLICANT: GRAY, Howard
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
TITLE OF INVENTION: CTL IMMUNITY
NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Kourie and Crew
STREET: Stewart Street Tower, One Market Plaza
CITY: San Francisco
STATE: California
COUNTRY: US
ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/197,484
FILING DATE: 16-FEB-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935,811
FILING DATE: 26-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,682
FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W.
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 467-9600
TELEFAX: (206) 623-6793
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide

US-08-197-484-72

Query Match 68.9%; Score 31; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 10;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTHDII 9
DB 1 TTHDII 6

RESULT 38
US-09-601-729-273
Sequence 273, Application US/09601729
Patent No. 6683052
GENERAL INFORMATION:
APPLICANT: THIAM, KADER
APPLICANT: AURIAULT, CLAUDE
APPLICANT: GRAS-MASSE, HELENE
APPLICANT: LOING, ESTELLE
APPLICANT: VERMAERDE, CLAUDE
APPLICANT: GUILLET, JEAN GERARD
TITLE OF INVENTION: LIPOPEPTIDES CONTAINING AN INTERFERON FRAGMENT AND USES
FILE REFERENCE: USB-97-AU-IN
CURRENT APPLICATION NUMBER: US/09/601,729
PRIOR FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: PCT/FR99/00259
PRIOR FILING DATE: 1999-02-05
PRIOR APPLICATION NUMBER: 98 01439
NUMBER OF SEQ ID NOS: 261
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 273
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-601-729-273

Query Match 68.9%; Score 31; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 10;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTHDII 9
DB 1 TTHDII 6

RESULT 39
PCT-US95-02121-72
Sequence 72, Application PCT/US9502121
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
TITLE OF INVENTION: CTL IMMUNITY
NUMBER OF SEQUENCES: 153
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/02121
FILING DATE: 16-FEB-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/197,484
FILING DATE: 16-FEB-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935,811

FILING DATE: 26-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,682
FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W.
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4PC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 467-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
PCT-US95-02121-72

Query Match 68.9%; Score 31; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 10;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTII 9
Db 1 TTTTII 6

RESULT 40
US-08-936-165A-297
Sequence 297, Application US/08936165A
Patent No. 6348582
GENERAL INFORMATION:
APPLICANT: Black, Michael
APPLICANT: Burnham, Martin
APPLICANT: Hodgson, John
APPLICANT: Knowles, David
APPLICANT: Lonetto, Michael
APPLICANT: Nicholas, Richard
APPLICANT: Pratt, Julie
APPLICANT: Reichard, Richard
APPLICANT: Rosenberg, Martin
APPLICANT: Ward, Judith
TITLE OF INVENTION: No. 6348582el Prokaryotic Polynucleotides,
TITLE OF INVENTION: Polypeptides and Their Uses
NUMBER OF SEQUENCES: 534
CORRESPONDENCE ADDRESS:
ADDRESSER: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
STATE: PA
COUNTRY: USA
ZIP: 19406-0939
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: Fast-Seq for Windows Version 2.0
CURRENT APPLICATION DATA:
FILING DATE: 24-SEP-1997
APPLICATION NUMBER: US/08/936,165A
CLASSIFICATION: 536
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/027,032
FILING DATE: 24-SEP-1996
ATTORNEY/AGENT INFORMATION:

NAME: Gimmi, Edward R.
REGISTRATION NUMBER: 38,891
REFERENCE/DOCKET NUMBER: B50549
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-4478
TELEFAX: 610-270-5090
TELEX:
INFORMATION FOR SEQ ID NO: 297:
SEQUENCE CHARACTERISTICS:
LENGTH: 56 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: Protein
US-08-936-165A-297

Query Match 68.9%; Score 31; DB 2; Length 56;
Best Local Similarity 55.6%; Pred. No. 61;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 LOTTTHDII 9
Db 7 MKRTTHDIL 15

RESULT 41
US-08-118-270-256
Sequence 256, Application US/08118270
Patent No. 5508384
GENERAL INFORMATION:
APPLICANT: Murphy, Randall B.
APPLICANT: Schuster, David I.
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
ADDRESSER: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,270
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033
REFERENCE/DOCKET NUMBER: MURPHY=2A
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO: 256:
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-118-270-256

Query Match 68.9%; Score 31; DB 1; Length 59;
Best Local Similarity 44.4%; Pred. No. 64;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

OY 1 LOTTHDII 9
|:|:|:|:
Db 26 LSSTHDLV 34

RESULT 42
US-09-390-027-6
Sequence 6, Application US/09390027
Patent No. 6235523
GENERAL INFORMATION:
APPLICANT: GATENCZYK, Diane M.
APPLICANT: PERSSON, Roy
APPLICANT: YAO, Fei-Long
APPLICANT: CAO, Shi-Xian
APPLICANT: KLEIN, Michel H.
APPLICANT: TARTAGLIA, James
APPLICANT: MOINGEON, Philippe
APPLICANT: ROVINSKI, Benjamin
TITLE OF INVENTION: TREATMENT OF CERVICAL CANCER
FILE REFERENCE: 1038-982 MIS-Jb
CURRENT APPLICATION NUMBER: US/09/390,027
CURRENT FILING DATE: 1999-09-03
EARLIER APPLICATION NUMBER: 60/099,291
EARLIER FILING DATE: 1998-09-04
NUMBER OF SEQ ID NOS: 12
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 59
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-390-027-6

Query Match 68.9%; Score 31; DB 2; Length 59;
Best Local Similarity 100.0%; Pred. No. 64;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
OY 4 TTHDII 9
|:|:|:|:
Db 50 TTHDII 55

RESULT 43
PCT-US93-08528-256
Sequence 256, Application PC/TUS9308528
GENERAL INFORMATION:
APPLICANT: New York University
TITLE OF INVENTION: POLYPEPTIDES OF G-COUPLED PROTEIN
TITLE OF INVENTION: RECEPTORS, AND COMPOSITIONS AND METHODS THEREOF
NUMBER OF SEQUENCES: 348
CORRESPONDENCE ADDRESS:
ADDRESSEE: BROWDY AND NEIMARK
STREET: 419 Seventh Street, N.W., Suite 300
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08528
FILING DATE: 09-SEP-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/943,236
FILING DATE: 10-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Townsend, Kevin G.
REGISTRATION NUMBER: 34,033

REFERENCE/DOCKET NUMBER: MURPHY-2 PCT
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-628-5197
TELEFAX: 202-737-3528
TELEX: 248633
INFORMATION FOR SEQ ID NO: 256:
SEQUENCE CHARACTERISTICS:
LENGTH: 59 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
PCT-US93-08528-256

Query Match 68.9%; Score 31; DB 4; Length 59;
Best Local Similarity 44.4%; Pred. No. 64;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

OY 1 LOTTHDII 9
|:|:|:|:
Db 26 LSSTHDLV 34

RESULT 44
US-08-785-795-1
Sequence 1, Application US/08785795
Patent No. 5932432
GENERAL INFORMATION:
APPLICANT: Ding N. Crowell, Branda J. Biermann
APPLICANT: and Stephen K. Randall
TITLE OF INVENTION: EXPRESSION LIBRARY SCREEN
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: Thomas Q. Henry
STREET: Bank One Center/Tower, Suite 3700
CITY: Indianapolis
STATE: Indiana
COUNTRY: USA
ZIP: 46204
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch,
MEDIUM TYPE: 1.4 Mb storage
COMPUTER: COMPAQ
OPERATING SYSTEM: MSDOS
SOFTWARE: ASCIT
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/785,795
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/283,875
FILING DATE: August 1, 1994
ATTORNEY/AGENT INFORMATION:
NAME: Thomas Q. Henry
REGISTRATION NUMBER: 28,309
REFERENCE/DOCKET NUMBER: IU-44
TELECOMMUNICATION INFORMATION:
TELEPHONE: (317) 634-3456
TELEFAX: (317) 637-7561
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 86 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-785-795-1

Query Match 68.9%; Score 31; DB 1; Length 86;
Best Local Similarity 57.1%; Pred. No. 96;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

OY 2 QTTHDI 8

Db 45 ETTLADV 51

RESULT 45
US-09-248-796A-21807
; Sequence 21807, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; PRIOR FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 21807
; LENGTH: 277
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-21807

Query Match 68.9%; Score 31; DB 2; Length 277;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 TTTTII 9
Db 95 TTTTII 100

RESULT 46
US-09-252-991A-27158
; Sequence 27158, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; PRIOR FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 27158
; LENGTH: 326
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-27158

Query Match 68.9%; Score 31; DB 2; Length 326;
Best Local Similarity 62.5%; Pred. No. 3.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 LQTTIHDI 8
Db 230 MTTTFHDI 237

RESULT 47
US-09-603-208A-62
; Sequence 62, Application US/09603208A
; Patent No. 6822084
; GENERAL INFORMATION:
; APPLICANT: Pompejus, Markus
; APPLICANT: Krieger, Burkhard

APPLICANT: Schroder, Hartwig
APPLICANT: Zelder, Oskar
APPLICANT: Habermann, Gregor
APPLICANT: Lee, Heung-Shick
APPLICANT: Kim, Hyung-Joon
TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING STRESS,
TITLE OF INVENTION: RESISTANCE AND TOLERANCE PROTEINS
FILE REFERENCE: BGI-124CP
CURRENT APPLICATION NUMBER: US/09/603,208A
CURRENT FILING DATE: 2000-06-23
PRIOR APPLICATION NUMBER: 60/141031
PRIOR FILING DATE: 1999-06-25
PRIOR APPLICATION NUMBER: 60/142692
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: 60/151214
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: DE 19930429.7
PRIOR FILING DATE: 1999-07-01
PRIOR APPLICATION NUMBER: DE 19931413.6
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19931457.8
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19931541.8
PRIOR FILING DATE: 1999-07-08
PRIOR APPLICATION NUMBER: DE 19932209.0
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: DE 19932230.9
PRIOR FILING DATE: 1999-07-09
PRIOR APPLICATION NUMBER: DE 19932914.1
PRIOR FILING DATE: 1999-07-14
PRIOR APPLICATION NUMBER: DE 19940764.9
PRIOR FILING DATE: 1999-08-27
PRIOR APPLICATION NUMBER: DE 19941382.7
PRIOR FILING DATE: 1999-08-31
NUMBER OF SEQ ID NOS: 306
SEQ ID NO 62
LENGTH: 383
TYPE: PRT
ORGANISM: Corynebacterium glutamicum
US-09-603-208A-62

Query Match 68.9%; Score 31; DB 2; Length 383;
Best Local Similarity 55.6%; Pred. No. 4.6e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 238 MKTTIHDI 246

RESULT 48
US-09-248-796A-20895
; Sequence 20895, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; PRIOR FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 20895
; LENGTH: 695
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-20895

Query Match 68.9%; Score 31; DB 2; Length 695;

Job time : 24.7 secs

Best Local Similarity 55.6%; Pred. No. 8.6e+02;
 Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
 :|||: ||:
 Db 29 VQTVEDIL 37

RESULT 49
 US-09-949-016-11304
 ; Sequence 11304, Application US/09949016
 ; Patent No. 6812339
 ; GENERAL INFORMATION:
 ; APPLICANT: VENTER, J. Craig et al.
 ; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
 ; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
 ; FILE REFERENCE: CL001307
 ; CURRENT APPLICATION NUMBER: US/09/949,016
 ; CURRENT FILING DATE: 2000-04-14
 ; PRIOR APPLICATION NUMBER: 60/241,755
 ; PRIOR FILING DATE: 2000-10-20
 ; PRIOR APPLICATION NUMBER: 60/237,768
 ; PRIOR FILING DATE: 2000-10-03
 ; PRIOR APPLICATION NUMBER: 60/231,498
 ; PRIOR FILING DATE: 2000-09-08
 ; NUMBER OF SEQ ID NOS: 207012
 ; SOFTWARE: FastSeq for Windows Version 4.0
 ; SEQ ID NO 11304
 ; LENGTH: 1016
 ; TYPE: PRT
 ; ORGANISM: Human
 ; US-09-949-016-11304

Query Match 68.9%; Score 31; DB 2; Length 1016;
 Best Local Similarity 62.5%; Pred. No. 1.3e+03;
 Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
 |||: ||:
 Db 368 LQRTVHDM 375

RESULT 50
 US-09-227-725A-4
 ; Sequence 4, Application US/09227725A
 ; Patent No. 6383758
 ; GENERAL INFORMATION:
 ; APPLICANT: St. George-Hylop, Peter H.
 ; APPLICANT: Rommens, Johanna
 ; APPLICANT: Fraser, Paul E.
 ; TITLE OF INVENTION: Alzheimer's Related Proteins and Methods
 ; TITLE OF INVENTION: of Use
 ; FILE REFERENCE: 1034/1P810-US1
 ; CURRENT APPLICATION NUMBER: US/09/227,725A
 ; CURRENT FILING DATE: 1999-01-08
 ; NUMBER OF SEQ ID NOS: 4
 ; SOFTWARE: FastSeq for Windows Version 3.0
 ; SEQ ID NO 4
 ; LENGTH: 1193
 ; TYPE: PRT
 ; ORGANISM: Homo Sapien
 ; US-09-227-725A-4

Query Match 68.9%; Score 31; DB 2; Length 1193;
 Best Local Similarity 62.5%; Pred. No. 1.5e+03;
 Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
 |||: ||:
 Db 355 LQRTVHDM 362

Search completed: May 5, 2006, 03:13:10

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OM protein - protein search, using sw model

Run on: May 5, 2006, 07:56:48 ; Search time 56 Seconds
(without alignments)
67.151 Million cell updates/sec

Title: US-08-170-344-6
Perfect score: 45
Sequence: 1 LQTTIHDI I 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Published Applications AA Main:*

- 1: /cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.rep:*
- 2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.rep:*
- 3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.rep:*
- 4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.rep:*
- 5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.rep:*
- 6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.rep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARYS

Result No.	Score	Query Match	Length	DB ID	Description
1	45	100.0	9	3	US-09-909-460-102 Sequence 102, App
2	45	100.0	9	3	US-09-872-835-102 Sequence 102, App
3	45	100.0	9	4	US-10-133-210-279 Sequence 279, App
4	45	100.0	9	5	US-10-758-970-102 Sequence 102, App
5	45	100.0	9	5	US-10-751-845-56 Sequence 56, App
6	45	100.0	10	5	US-10-484-063-22 Sequence 22, App
7	45	100.0	15	4	US-10-476-570-21 Sequence 21, App
8	45	100.0	15	4	US-10-476-570-22 Sequence 22, App
9	45	100.0	21	4	US-10-476-570-8 Sequence 8, App
10	45	100.0	30	5	US-10-476-570-53 Sequence 53, App
11	45	100.0	30	5	US-10-858-384-4 Sequence 4, App
12	45	100.0	32	4	US-10-476-570-9 Sequence 9, App
13	45	100.0	33	4	US-10-476-570-19 Sequence 19, App
14	45	100.0	151	4	US-10-177-390-6 Sequence 6, App
15	45	100.0	151	5	US-10-484-063-20 Sequence 20, App
16	45	100.0	151	5	US-10-484-063-27 Sequence 27, App
17	45	100.0	158	5	US-10-858-384-2 Sequence 2, App
18	45	100.0	158	5	US-10-367-057-16 Sequence 16, App
19	45	100.0	158	6	US-11-021-949-13 Sequence 13, App
20	45	100.0	171	4	US-10-472-724-2 Sequence 2, App
21	45	100.0	243	6	US-11-072-288-1 Sequence 1, App
22	45	100.0	266	3	US-09-367-309A-1 Sequence 1, App
23	45	100.0	273	4	US-10-000-903-4 Sequence 4, App
24	45	100.0	273	4	US-10-899-771-4 Sequence 4, App
25	45	100.0	292	4	US-10-000-903-10 Sequence 10, App
26	45	100.0	292	5	US-10-899-771-10 Sequence 10, App
27	45	100.0	371	4	US-10-000-903-6 Sequence 6, App

28	45	100.0	371	5	US-10-899-771-6 Sequence 6, App
29	45	100.0	390	4	US-10-000-903-14 Sequence 14, App
30	45	100.0	390	5	US-10-899-771-14 Sequence 14, App
31	45	100.0	536	4	US-10-367-095-10 Sequence 10, App
32	45	100.0	536	4	US-10-368-046-10 Sequence 10, App
33	45	100.0	536	4	US-10-367-367-10 Sequence 10, App
34	45	100.0	536	4	US-10-918-337-10 Sequence 10, App
35	41	91.1	25	6	US-11-021-949-1 Sequence 1, App
36	37	82.2	248	5	US-10-820-155-42 Sequence 42, App
37	37	82.2	248	5	US-10-820-155-57 Sequence 57, App
38	36	80.0	15	4	US-10-476-570-23 Sequence 23, App
39	36	80.0	151	4	US-10-425-115-237192 Sequence 237192, App
40	36	80.0	601	6	US-11-097-143-4221 Sequence 4221, App
41	36	80.0	1082	4	US-10-282-122A-49664 Sequence 49664, App
42	36	80.0	1198	4	US-10-452-024-85 Sequence 85, App
43	36	80.0	1381	6	US-11-097-143-36795 Sequence 36795, App
44	35	77.8	97	4	US-10-767-701-50864 Sequence 50864, App
45	35	77.8	156	4	US-10-425-115-263392 Sequence 263392, App
46	35	77.8	165	4	US-10-425-114-55470 Sequence 55470, App
47	35	77.8	316	4	US-10-424-599-168462 Sequence 168462, App
48	35	77.8	336	4	US-10-424-599-230650 Sequence 230650, App
49	35	77.8	403	4	US-10-437-963-117225 Sequence 117225, App
50	35	77.8	448	4	US-10-425-115-263390 Sequence 263390, App
51	35	77.8	463	4	US-10-425-114-49312 Sequence 49312, App
52	35	77.8	537	4	US-10-424-599-230650 Sequence 230650, App
53	35	77.8	540	4	US-10-424-599-230656 Sequence 230656, App
54	35	77.8	552	4	US-10-425-114-46247 Sequence 46247, App
55	35	77.8	552	4	US-10-425-114-46829 Sequence 46829, App
56	34	75.6	158	6	US-11-021-949-361 Sequence 361, App
57	34	75.6	233	4	US-10-772-212A-16 Sequence 16, App
58	34	75.6	619	4	US-10-320-797-3259 Sequence 3259, App
59	33	73.3	40	4	US-10-425-115-119095 Sequence 319095, App
60	33	73.3	59	4	US-10-425-115-224781 Sequence 224781, App
61	33	73.3	102	4	US-10-425-115-168030 Sequence 168030, App
62	33	73.3	149	6	US-11-021-949-360 Sequence 360, App
63	33	73.3	162	6	US-11-021-949-31 Sequence 31, App
64	33	73.3	193	4	US-10-017-161-46 Sequence 46, App
65	33	73.3	193	4	US-10-450-763-53459 Sequence 53459, App
66	33	73.3	198	5	US-10-408-765A-2849 Sequence 2849, App
67	33	73.3	297	4	US-10-288-160-8 Sequence 8, App
68	33	73.3	302	5	US-10-774-355A-2386 Sequence 2386, App
69	33	73.3	310	5	US-10-174-355A-1784 Sequence 1784, App
70	33	73.3	323	4	US-10-188-186-18 Sequence 18, App
71	33	73.3	340	4	US-10-250-615-18 Sequence 18, App
72	33	73.3	496	4	US-10-424-599-155954 Sequence 155954, App
73	33	73.3	500	4	US-10-214-269-19 Sequence 19, App
74	33	73.3	501	5	US-10-733-923-9631 Sequence 9631, App
75	33	73.3	501	5	US-10-733-923-9692 Sequence 9692, App
76	33	73.3	568	6	US-11-097-143-28185 Sequence 28185, App
77	33	73.3	779	6	US-11-097-143-150 Sequence 150, App
78	33	73.3	839	4	US-10-399-456-11 Sequence 11, App
79	33	73.3	53	4	US-10-767-701-54673 Sequence 54673, App
80	32	71.1	58	4	US-10-424-599-255207 Sequence 255207, App
81	32	71.1	69	4	US-10-425-115-280425 Sequence 280425, App
82	32	71.1	117	4	US-10-437-963-128330 Sequence 128330, App
83	32	71.1	137	4	US-10-437-963-151775 Sequence 151775, App
84	32	71.1	147	4	US-10-437-963-161775 Sequence 161775, App
85	32	71.1	168	4	US-10-437-963-201751 Sequence 201751, App
86	32	71.1	168	4	US-10-424-599-197053 Sequence 197053, App
87	32	71.1	238	4	US-10-425-115-342845 Sequence 342845, App
88	32	71.1	241	4	US-10-369-493-10503 Sequence 10503, App
89	32	71.1	259	4	US-10-424-599-171760 Sequence 171760, App
90	32	71.1	333	4	US-10-369-493-10175 Sequence 10175, App
91	32	71.1	333	4	US-10-369-493-20899 Sequence 20899, App
92	32	71.1	404	6	US-11-097-143-13188 Sequence 13188, App
93	32	71.1	420	5	US-10-450-763-43256 Sequence 43256, App
94	32	71.1	420	5	US-10-450-763-45796 Sequence 45796, App
95	32	71.1	420	5	US-10-450-763-51741 Sequence 51741, App
96	32	71.1	481	6	US-11-097-143-8211 Sequence 8211, App
97	32	71.1	485	4	US-10-389-566-2343 Sequence 2343, App
98	32	71.1	497	4	US-10-369-435-18 Sequence 18, App
99	32	71.1	513	3	US-09-769-787-37 Sequence 37, App
100	32	71.1			

101	32	71.1	513	3	US-09-769-787-193	Sequence 193, App	174	31	68.9	687	5	US-10-370-715B-538	Sequence 538, App
102	32	71.1	513	3	US-10-472-928-2686	Sequence 2686, Ap	175	31	68.9	703	5	US-10-450-763-48057	Sequence 48057, A
103	32	71.1	553	4	US-10-467-534-61	Sequence 61, Appl	176	31	68.9	764	4	US-10-437-963-151490	Sequence 151490,
104	32	71.1	853	4	US-10-032-585-7830	Sequence 7830, Ap	177	31	68.9	785	4	US-10-264-049-2968	Sequence 2968, Ap
105	32	71.1	870	5	US-10-450-763-60646	Sequence 60646, A	178	31	68.9	785	5	US-10-450-763-51180	Sequence 51180, A
106	32	71.1	1104	4	US-10-369-493-22548	Sequence 22548, A	179	31	68.9	834	4	US-10-094-749-2227	Sequence 2227, Ap
107	32	71.1	1258	4	US-10-310-154-703	Sequence 703, App	180	31	68.9	841	4	US-10-467-490-2	Sequence 2, Appl1
108	32	71.1	1258	5	US-10-732-923-621	Sequence 621, App	181	31	68.9	877	3	US-09-934-070-10	Sequence 10, Appl
109	32	71.1	1258	5	US-10-732-923-621	Sequence 621, App	182	31	68.9	897	3	US-10-222-772-10	Sequence 33, Appl
110	32	71.1	1280	6	US-11-037-143-1599	Sequence 22696, A	183	31	68.9	901	3	US-09-737-149-33	Sequence 2, Appl1
111	32	71.1	1288	5	US-10-745-237-180	Sequence 1599, Ap	184	31	68.9	901	4	US-10-416-793-2	Sequence 33, Appl
112	32	71.1	1585	4	US-10-437-963-124349	Sequence 124349,	185	31	68.9	901	5	US-10-701-283-33	Sequence 4, Appl1
113	31	68.9	9	4	US-10-777-053-547	Sequence 547, App	186	31	68.9	908	5	US-10-493-047-9	Sequence 87, Appl
114	31	68.9	9	4	US-10-837-217-547	Sequence 27, App	187	31	68.9	908	5	US-10-805-684-87	Sequence 4, Appl1
115	31	68.9	10	4	US-10-128-711-72	Sequence 280, App	188	31	68.9	965	3	US-09-737-149-4	Sequence 4, Appl1
116	31	68.9	10	4	US-10-133-210-280	Sequence 20, Appl	189	31	68.9	965	3	US-10-701-283-4	Sequence 4, Appl1
117	31	68.9	15	4	US-10-476-570-20	Sequence 20, App	190	31	68.9	971	3	US-09-737-149-6	Sequence 6, Appl1
118	31	68.9	22	4	US-10-639-067-204	Sequence 204, App	191	31	68.9	971	4	US-10-701-283-6	Sequence 3, Appl1
119	31	68.9	48	4	US-10-424-599-153049	Sequence 153049,	192	31	68.9	977	4	US-10-467-490-3	Sequence 2626, Ap
120	31	68.9	48	4	US-10-437-963-157793	Sequence 157793,	193	31	68.9	995	4	US-10-094-749-2626	Sequence 2, Appl1
121	31	68.9	49	4	US-10-437-963-157793	Sequence 157793,	194	31	68.9	998	4	US-10-106-534-2	Sequence 6, Appl
122	31	68.9	53	4	US-10-424-599-258195	Sequence 258195,	195	31	68.9	1008	4	US-10-222-772-62	Sequence 6, Appl1
123	31	68.9	56	3	US-10-425-115-269612	Sequence 269612,	196	31	68.9	1011	3	US-09-934-070-6	Sequence 6, Appl1
124	31	68.9	68	4	US-09-939-980-297	Sequence 297, App	197	31	68.9	1011	4	US-10-222-772-6	Sequence 8, Appl1
125	31	68.9	68	4	US-10-437-963-163168	Sequence 163168,	198	31	68.9	1037	3	US-09-737-149-8	Sequence 8, Appl1
126	31	68.9	78	4	US-10-424-599-236566	Sequence 236566,	199	31	68.9	1043	4	US-10-701-283-8	Sequence 8, Appl1
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128	31	68.9	87	4	US-10-424-599-358669	Sequence 358669,	201	31	68.9	1061	4	US-10-437-963-197935	Sequence 197935, Ap
129	31	68.9	87	4	US-10-425-115-358669	Sequence 209999,	202	31	68.9	1130	4	US-10-282-122A-49890	Sequence 12, Appl
130	31	68.9	88	4	US-10-424-599-277308	Sequence 277308,	203	31	68.9	1130	4	US-10-149-819-12	Sequence 67, Appl
131	31	68.9	99	4	US-10-424-599-165390	Sequence 165390,	204	31	68.9	1193	3	US-10-260-708-67	Sequence 4, Appl1
132	31	68.9	104	4	US-10-425-115-304710	Sequence 304710,	205	31	68.9	1211	4	US-10-071-900-4	Sequence 4487, Ap
133	31	68.9	110	4	US-10-767-701-62097	Sequence 62097, A	206	31	68.9	1211	4	US-09-969-034-4487	Sequence 14, Appl
134	31	68.9	112	4	US-10-767-701-38587	Sequence 38587, A	207	31	68.9	1632	4	US-10-242-943-14	Sequence 14, Appl
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137	31	68.9	146	4	US-10-424-599-278042	Sequence 278042,	210	31	68.9	2042	4	US-10-450-763-38020	Sequence 38020, A
138	31	68.9	150	4	US-10-437-963-117374	Sequence 117374,	211	31	68.9	2042	4	US-10-192-584-6	Sequence 296, Appl1
139	31	68.9	166	4	US-10-425-115-277001	Sequence 277001,	212	31	68.9	2296	5	US-10-092-900A-295	Sequence 295, Ap
140	31	68.9	177	4	US-10-425-115-277001	Sequence 36396, A	213	31	68.9	2296	5	US-10-753-860-1555	Sequence 1555, Ap
141	31	68.9	180	6	US-11-097-143-36396	Sequence 32448, A	214	31	68.9	2327	4	US-10-766-149-5112	Sequence 5112, Ap
142	31	68.9	183	4	US-10-029-386-32448	Sequence 209499,	215	31	68.9	2385	4	US-10-092-900A-294	Sequence 294, App
143	31	68.9	187	4	US-10-425-115-209499	Sequence 21504, A	216	31	68.9	2390	4	US-10-282-122A-46587	Sequence 46587, A
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147	31	68.9	306	4	US-10-425-114-51732	Sequence 8092, Ap	220	30	66.7	54	4	US-10-425-115-320406	Sequence 320406,
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158	31	68.9	420	4	US-10-424-599-257607	Sequence 457608,	231	30	66.7	112	5	US-10-970-713-322	Sequence 322, App
159	31	68.9	420	4	US-10-424-599-257608	Sequence 46164, A	232	30	66.7	115	4	US-10-424-599-211982	Sequence 211982,
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274	30	66.7	247	4	US-10-425-114-52506	Sequence 52506, A	347	30	66.7	522	3	US-10-976-800-86	Sequence 96, Appl
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278	30	66.7	248	5	US-10-971-461-26	Sequence 26, Appl	351	30	66.7	522	4	US-10-139-296-96	Sequence 96, Appl
279	30	66.7	248	5	US-10-820-155-58	Sequence 58, Appl	352	30	66.7	522	4	US-10-139-296-96	Sequence 96, Appl
280	30	66.7	248	5	US-10-756-149-5337	Sequence 5337, Ap	353	30	66.7	522	4	US-10-139-218-86	Sequence 96, Appl
281	30	66.7	259	3	US-09-925-302-473	Sequence 473, App	354	30	66.7	522	4	US-10-405-660-86	Sequence 96, Appl
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403	30	66.7	1315	4	US-10-697-036-13	Sequence 13, Appl	476	29	64.4	208	4	US-10-335-977-9731	Sequence 9731, Ap
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406	30	66.7	1434	5	US-10-494-672-220	Sequence 220, App	479	29	64.4	215	5	US-10-869-630-26	Sequence 26, Appl
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409	30	66.7	1538	4	US-10-437-963-151793	Sequence 151793,	482	29	64.4	223	4	US-10-425-114-62545	Sequence 62545, A
410	30	66.7	1807	5	US-10-723-860-1336	Sequence 1336, Ap	483	29	64.4	227	4	US-10-424-599-206469	Sequence 206469,
411	30	66.7	2039	4	US-10-192-584-7	Sequence 7, Appl1	484	29	64.4	228	4	US-10-425-115-229884	Sequence 229884,
412	29	64.4	20	4	US-10-083-919A-27	Sequence 27, Appl1	485	29	64.4	229	4	US-10-282-122A-71091	Sequence 71091, A
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416	29	64.4	51	4	US-09-864-761-36345	Sequence 36345, A	489	29	64.4	237	4	US-10-724-972A-3781	Sequence 3781, Ap
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422	29	64.4	58	4	US-10-425-115-191848	Sequence 191848,	495	29	64.4	250	4	US-09-925-998-5	Sequence 508, App
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433	29	64.4	79	4	US-10-437-963-118212	Sequence 118212,	506	29	64.4	301	4	US-10-424-599-235633	Sequence 235633,
434	29	64.4	80	4	US-10-425-115-226367	Sequence 226367,	507	29	64.4	304	4	US-10-424-599-235630	Sequence 149433,
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438	29	64.4	86	4	US-10-083-919A-100	Sequence 100, App	511	29	64.4	311	4	US-10-032-201B-216	Sequence 6561, Ap
439	29	64.4	86	4	US-10-083-919A-183	Sequence 183, App	512	29	64.4	312	5	US-10-739-930-6561	Sequence 11099, A
440	29	64.4	90	4	US-10-106-698-5200	Sequence 6200, Ap	513	29	64.4	334	4	US-10-156-761-11099	Sequence 249040,
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443	29	64.4	99	5	US-10-739-930-6347	Sequence 6347, Ap	516	29	64.4	332	5	US-10-276-825-13	Sequence 39744, A
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446	29	64.4	103	4	US-10-437-963-145205	Sequence 145205,	519	29	64.4	335	4	US-10-369-493-21664	Sequence 486, App
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453	29	64.4	129	3	US-09-928-298-443	Sequence 443, App	526	29	64.4	341	4	US-10-424-599-155318	Sequence 155318,
454	29	64.4	129	4	US-10-102-806-443	Sequence 443, App	527	29	64.4	345	4	US-10-085-198-130	Sequence 130, App
455	29	64.4	129	4	US-10-425-115-336015	Sequence 236015,	528	29	64.4	345	4	US-10-000-012-20	Sequence 49993, A
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461	29	64.4	145	6	US-11-097-143-39516	Sequence 39516, A	534	29	64.4	352	4	US-10-425-115-224369	Sequence 23280, A
462	29	64.4	149	6	US-11-021-949-15	Sequence 15, Appl	535	29	64.4	354	4	US-10-424-599-21280	Sequence 56070, A
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543	29	64.4	395	4	US-10-697-828-7	Sequence 7, Appl	616	29	64.4	628	6	US-11-051-454-446	Sequence 246, App
544	29	64.4	395	4	US-10-841-707-8	Sequence 8, Appl	617	29	64.4	634	6	US-11-067-557-10	Sequence 10, App
545	29	64.4	408	4	US-10-214-446-4	Sequence 4, Appl	618	29	64.4	636	5	US-10-450-763-35986	Sequence 35986, A
546	29	64.4	411	4	US-10-282-122A-61391	Sequence 61391, A	619	29	64.4	650	4	US-10-369-493-6555	Sequence 2655, App
547	29	64.4	415	4	US-10-282-122A-70442	Sequence 70442, A	620	29	64.4	656	5	US-10-998-197-82	Sequence 52, Appl
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549	29	64.4	418	3	US-09-927-602-5	Sequence 5, Appl	622	29	64.4	684	3	US-09-988-626-333	Sequence 233, App
550	29	64.4	426	4	US-10-282-122A-51811	Sequence 51811, A	623	29	64.4	684	3	US-09-988-687-233	Sequence 233, App
551	29	64.4	431	4	US-10-012-819-228	Sequence 228, App	624	29	64.4	684	3	US-09-988-686-233	Sequence 20, Appl
552	29	64.4	431	5	US-10-739-930-11039	Sequence 11039, A	625	29	64.4	696	5	US-10-480-988-25	Sequence 25, Appl
553	29	64.4	432	5	US-10-739-930-11041	Sequence 11041, A	626	29	64.4	701	4	US-10-237-895A-19	Sequence 19, Appl
554	29	64.4	433	2	US-08-945-038-6	Sequence 6, Appl	627	29	64.4	714	5	US-10-276-825-12	Sequence 12, Appl
555	29	64.4	433	4	US-10-335-977-8500	Sequence 8500, Ap	628	29	64.4	715	5	US-10-276-825-15	Sequence 15, Appl
556	29	64.4	436	4	US-10-094-886-152	Sequence 152, App	629	29	64.4	726	4	US-10-424-599-262675	Sequence 262675,
557	29	64.4	436	4	US-10-732-923-8388	Sequence 8388, Ap	630	29	64.4	728	4	US-10-437-963-129070	Sequence 129070,
558	29	64.4	437	3	US-09-973-322-5	Sequence 11, Appl	631	29	64.4	737	4	US-10-106-698-4893	Sequence 4893, Ap
559	29	64.4	437	3	US-10-170-385-313	Sequence 11, Appl	632	29	64.4	738	4	US-10-087-132-762	Sequence 762, App
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561	29	64.4	437	5	US-10-417-375-50	Sequence 5, Appl	634	29	64.4	738	5	US-10-723-860-2006	Sequence 2006, Ap
562	29	64.4	437	5	US-10-417-375-52	Sequence 52, Appl	635	29	64.4	738	5	US-10-276-825-11	Sequence 11, Appl
563	29	64.4	437	5	US-10-417-375-54	Sequence 54, Appl	636	29	64.4	738	5	US-10-276-825-18	Sequence 18, Appl
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565	29	64.4	438	4	US-10-335-977-8501	Sequence 5501, Ap	638	29	64.4	738	5	US-10-287-436A-1160	Sequence 1160, Ap
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567	29	64.4	444	3	US-09-815-242-11091	Sequence 11091, A	640	29	64.4	761	4	US-10-225-486-55	Sequence 55, Appl
568	29	64.4	447	3	US-09-835-684-5	Sequence 5, Appl	641	29	64.4	761	4	US-10-408-765A-696	Sequence 696, App
569	29	64.4	447	3	US-09-880-371-5	Sequence 5, Appl	642	29	64.4	761	4	US-10-723-860-3785	Sequence 3785, App
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571	29	64.4	447	4	US-10-010-390-5	Sequence 5, Appl	644	29	64.4	764	5	US-10-369-493-6493	Sequence 6493, Ap
572	29	64.4	447	4	US-10-441-736-6	Sequence 6, Appl	645	29	64.4	766	4	US-10-297-895A-21	Sequence 21, Appl
573	29	64.4	447	5	US-10-847-142-5	Sequence 5, Appl	646	29	64.4	804	4	US-10-437-963-303648	Sequence 303648,
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585	29	64.4	486	4	US-10-354-358-64	Sequence 64, Appl	658	29	64.4	1001	4	US-10-438-753-4	Sequence 4, Appl
586	29	64.4	496	4	US-10-425-114-45726	Sequence 45726, A	659	29	64.4	1001	4	US-10-438-753-5	Sequence 5, Appl
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591	29	64.4	496	5	US-10-756-149-5314	Sequence 5314, Ap	664	29	64.4	1053	4	US-10-041-018-203	Sequence 203, App
592	29	64.4	502	4	US-10-214-263-16	Sequence 16, Appl	665	29	64.4	1053	4	US-10-041-018-227	Sequence 227, App
593	29	64.4	502	4	US-10-437-963-198057	Sequence 198057,	666	29	64.4	1053	4	US-10-041-018-227	Sequence 287, App
594	29	64.4	509	5	US-10-732-923-1145	Sequence 1145, Ap	667	29	64.4	1053	4	US-10-041-018-339	Sequence 339, App
595	29	64.4	516	4	US-10-156-761-10185	Sequence 10185, A	668	29	64.4	1137	4	US-10-437-963-177112	Sequence 177112,
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597	29	64.4	527	4	US-10-078-929-106	Sequence 106, App	670	29	64.4	1245	4	US-10-733-923-8336	Sequence 8336, Ap
598	29	64.4	527	4	US-10-102-806-490	Sequence 490, App	671	29	64.4	1268	4	US-10-437-963-811037	Sequence 811037,
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601	29	64.4	553	4	US-10-425-115-278342	Sequence 278342,	674	29	64.4	1688	4	US-10-087-168-82	Sequence 82, Appl
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604	29	64.4	586	4	US-10-369-493-6115	Sequence 6115, Ap	677	29	64.4	1894	4	US-10-218-779-81	Sequence 81, Appl
605	29	64.4	590	5	US-10-631-467-1494	Sequence 1494, Ap	678	29	64.4	1894	4	US-10-451-010-9	Sequence 9, Appl
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607	29	64.4	600	4	US-10-282-122A-69214	Sequence 69214, A	680	29	64.4	1896	3	US-10-312-352-34	Sequence 34, Appl
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609	29	64.4	613	4	US-10-458-143-14	Sequence 14, Appl	682	29	64.4	1925	4	US-10-087-684-32	Sequence 32, Appl
610	29	64.4	623	5	US-10-739-930-9175	Sequence 9175, Ap	683	29	64.4	1925	4	US-10-087-684-32	Sequence 32, Appl
611	29	64.4	628	4	US-10-176-847-108	Sequence 108, App	684	29	64.4	1925	4	US-10-218-779-32	Sequence 32, Appl

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686	29	64.4	2110	5	US-10-732-923-8666	Sequence 8666, Ap	759	28	62.2	102	2	US-08-260-675-12	Sequence 12, Appl
687	29	64.4	2210	5	US-10-741-600-1318	Sequence 1218, Ap	760	28	62.2	102	3	US-09-952-118A-12	Sequence 12, Appl
688	29	64.4	2211	5	US-10-741-600-1324	Sequence 1324, Ap	761	28	62.2	102	3	US-09-754-831A-17	Sequence 7, Appl
689	29	64.4	2244	5	US-10-741-600-1321	Sequence 1321, Ap	762	28	62.2	102	4	US-10-050-050-12	Sequence 12, Appl
690	29	64.4	2411	5	US-10-741-600-1323	Sequence 1323, Ap	763	28	62.2	102	4	US-10-187-394-7	Sequence 7, Appl
691	29	64.4	2887	5	US-10-479-875-8	Sequence 8, Appl	764	28	62.2	102	4	US-10-187-394-8	Sequence 8, Appl
692	29	64.4	2977	5	US-10-741-600-1320	Sequence 1320, Ap	765	28	62.2	102	4	US-10-424-599-230042	Sequence 230042,
693	29	64.4	3151	5	US-10-486-678-16	Sequence 16, Appl	766	28	62.2	102	4	US-10-385-064-12	Sequence 12, Appl
694	29	64.4	3173	3	US-09-918-715-218	Sequence 218, App	767	28	62.2	102	5	US-10-671-217-7	Sequence 7, Appl
695	29	64.4	3173	3	US-10-474-794-218	Sequence 218, App	768	28	62.2	102	5	US-10-865-514-12	Sequence 12, Appl
696	29	64.4	3173	5	US-10-979-159-218	Sequence 218, App	769	28	62.2	102	6	US-11-037-782-165	Sequence 165, App
697	29	64.4	3176	4	US-10-372-683-10	Sequence 20, Appl	770	28	62.2	106	4	US-10-425-115-264146	Sequence 264146,
698	29	64.4	3176	4	US-10-734-864-91	Sequence 91, Appl	771	28	62.2	106	4	US-10-425-115-271754	Sequence 271754,
699	29	64.4	3176	5	US-10-723-860-1065	Sequence 1065, Ap	772	28	62.2	107	4	US-10-425-115-371771	Sequence 371771,
700	29	64.4	3176	5	US-10-486-678-15	Sequence 15, Appl	773	28	62.2	110	4	US-10-425-115-371771	Sequence 371771,
701	29	64.4	3176	5	US-10-853-335A-160	Sequence 160, App	774	28	62.2	117	4	US-10-437-963-122278	Sequence 122278,
702	29	64.4	3176	5	US-10-287-436A-503	Sequence 503, App	775	28	62.2	118	3	US-09-389-705-11	Sequence 49077, A
703	29	64.4	3176	5	US-10-287-436A-1196	Sequence 1196, Ap	776	28	62.2	118	4	US-10-115-406-8	Sequence 10818, A
704	29	64.4	3177	5	US-10-741-600-1319	Sequence 1319, Ap	777	28	62.2	118	4	US-10-154-333-10	Sequence 6, Appl
705	29	64.4	3494	5	US-10-473-127-697	Sequence 697, App	778	28	62.2	118	4	US-10-704-223-8	Sequence 8, Appl
706	29	64.4	3494	5	US-10-473-127-697	Sequence 703, App	779	28	62.2	120	4	US-10-425-115-362046	Sequence 362046,
707	29	64.4	3623	5	US-10-741-601-335	Sequence 703, App	780	28	62.2	126	4	US-10-767-701-50300	Sequence 50300, A
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710	29	64.4	3623	5	US-10-473-127-702	Sequence 702, App	783	28	62.2	134	4	US-10-767-701-31594	Sequence 31594, A
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713	29	64.4	3631	5	US-10-282-122A-71235	Sequence 71235, A	786	28	62.2	137	4	US-10-425-115-313156	Sequence 10818, A
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715	28.5	63.3	4640	5	US-10-408-765A-2103	Sequence 2103, Ap	788	28	62.2	142	4	US-10-424-599-166248	Sequence 6, Appl
716	28.5	63.3	4641	5	US-10-850-581-2	Sequence 2, Appl	789	28	62.2	144	4	US-10-002-478-6	Sequence 210797,
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721	28	62.2	31	4	US-10-643-627-21	Sequence 21, Appl	794	28	62.2	153	5	US-10-732-923-8264	Sequence 48367, A
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727	28	62.2	60	4	US-10-424-599-375730	Sequence 275730,	800	28	62.2	171	3	US-09-927-602-8	Sequence 210797,
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752	28	62.2	94	5	US-10-733-930-9180	Sequence 9180, Ap	825	28	62.2	212	4	US-10-158-057-260	Sequence 260, App
753	28	62.2	97	4	US-10-424-599-150518	Sequence 150518,	826	28	62.2	216	4	US-10-425-114-466138	Sequence 466138,
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832	28	62.2	241	3	US-09-738-626-6424	Sequence 6424, App	905	28	62.2	256	4	US-10-183-010-374	Sequence 374, App
833	28	62.2	241	5	US-10-739-930-8715	Sequence 8715, App	906	28	62.2	256	4	US-10-183-012-374	Sequence 374, App
834	28	62.2	241	5	US-10-347-669-32	Sequence 32, App1	907	28	62.2	256	4	US-10-184-614-374	Sequence 374, App
835	28	62.2	241	6	US-11-006-098-50	Sequence 50, App1	908	28	62.2	256	4	US-10-184-623-374	Sequence 374, App
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838	28	62.2	247	5	US-10-820-155-78	Sequence 78, App1	911	28	62.2	256	4	US-10-184-646-374	Sequence 374, App
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846	28	62.2	256	4	US-10-174-590-374	Sequence 374, App	919	28	62.2	256	4	US-10-196-464-374	Sequence 374, App
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848	28	62.2	256	4	US-10-175-737-374	Sequence 374, App	921	28	62.2	256	4	US-10-176-751-374	Sequence 374, App
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857	28	62.2	256	4	US-10-176-482-374	Sequence 374, App	930	28	62.2	256	4	US-10-183-013-374	Sequence 374, App
858	28	62.2	256	4	US-10-176-757-374	Sequence 374, App	931	28	62.2	256	4	US-10-184-612-374	Sequence 37

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ALIGNMENTS

RESULT 1
US-09-909-460-102
; Sequence 102, Application US/09909460
; Publication No. US20020182258A1
; GENERAL INFORMATION:
; APPLICANT: Lumsford, Lynn B.
; APPLICANT: Putnam, David
; APPLICANT: Hedley, Mary Lynn
; TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY OF NUCLEIC
; FILE REFERENCE: 08191/014001
; CURRENT APPLICATION NUMBER: US/09/909,460
; PRIOR FILING DATE: 2001-07-18
; PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: US/09/321,346
; NUMBER OF SEQ ID NOS: 114
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 102
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Hepatitis B virus
US-09-909-460-102

Query Match 100.0%; Score 45; DB 3; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 1 LQTTIHDI 9

RESULT 2
US-09-872-836-102
; Sequence 102, Application US/09872836
; Publication No. US20040142475A1
; GENERAL INFORMATION:
; APPLICANT: Barman, Shikha P.
; APPLICANT: McKeever, Una
; APPLICANT: Hedley, Mary Lynne
; TITLE OF INVENTION: DELIVERY SYSTEMS FOR BIOACTIVE AGENTS
; FILE REFERENCE: 08191-018001
; CURRENT APPLICATION NUMBER: US/09/872,836
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 102

; PRIOR APPLICATION NUMBER: US 60/208,830
; PRIOR FILING DATE: 2000-06-02
; NUMBER OF SEQ ID NOS: 120
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 102
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Homo sapiens
US-09-872-836-102

Query Match 100.0%; Score 45; DB 3; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 1 LQTTIHDI 9

RESULT 3
US-10-133-210-279
; Sequence 279, Application US/10133210
; Publication No. US20030103964A1
; GENERAL INFORMATION:
; APPLICANT: Berlinsky, Jay
; APPLICANT: Gulukota, Kamalakara
; APPLICANT: Vaccaro, Dennis
; APPLICANT: Wang, Zhiping
; APPLICANT: Zhang, Chao
; TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
; FILE REFERENCE: BU-035AX
; CURRENT APPLICATION NUMBER: US/10/133,210
; PRIOR FILING DATE: 2002-04-26
; NUMBER OF SEQ ID NOS: 281
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 279
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-279

Query Match 100.0%; Score 45; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 1 LQTTIHDI 9

RESULT 4
US-10-758-970-102
; Sequence 102, Application US/10758970
; Publication No. US20050037086A1
; GENERAL INFORMATION:
; APPLICANT: Hedley, Mary Lynne
; APPLICANT: Heu, Yung-Yueh
; APPLICANT: Iyo, Michael
; TITLE OF INVENTION: CONTINUOUS-FLOW METHOD FOR PREPARING MICROPARTICLES
; FILE REFERENCE: 08191-012001
; CURRENT APPLICATION NUMBER: US/10/758,970
; PRIOR FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: US/09/715,708A
; PRIOR FILING DATE: 2000-11-17
; PRIOR APPLICATION NUMBER: US 60/166,516
; NUMBER OF SEQ ID NOS: 109
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 102

LENGTH: 9
TYPE: PRT
ORGANISM: Human papilloma virus
US-10-758-970-102

Query Match 100.0%; Score 45; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LQTTIHDI 9
Db 1 LQTTIHDI 9

RESULT 5
US-10-751-845-56
Sequence 56, Application US/10751845
Publication No. US20050100928A1
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Urban, Robert G.
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING POLYPEPTIDE POLYPEPTIDES
FILE REFERENCE: 08191-013001
CURRENT FILING DATE: US/10/751,845
PRIOR APPLICATION NUMBER: US/09/664,225
PRIOR FILING DATE: 2004-01-05
PRIOR FILING DATE: 2000-08-18
PRIOR APPLICATION NUMBER: US 60/169,846
PRIOR FILING DATE: 1999-12-09
PRIOR APPLICATION NUMBER: US 60/154,665
PRIOR FILING DATE: 1999-09-16
NUMBER OF SEQ ID NOS: 163
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 56
LENGTH: 9
TYPE: PRT
ORGANISM: Human Papilloma virus
US-10-751-845-56

Query Match 100.0%; Score 45; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LQTTIHDI 9
Db 1 LQTTIHDI 9

RESULT 6
US-10-484-063-2
Sequence 2, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLEMO
APPLICANT: FOLEN, MICHAEL
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
PRIOR FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 2
LENGTH: 10
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-484-063-2

Query Match 100.0%; Score 45; DB 5; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LQTTIHDI 9
Db 2 LQTTIHDI 10

RESULT 7
US-10-476-570-21
Sequence 21, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from B6 and/or B7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 21
LENGTH: 15
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide B6 20-34
US-10-476-570-21

Query Match 100.0%; Score 45; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.09;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 LQTTIHDI 9
Db 7 LQTTIHDI 15

RESULT 8
US-10-476-570-22
Sequence 22, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from B6 and/or B7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 22
LENGTH: 15

;; TYPE: PRT
;; ORGANISM: artificial sequence
;; FEATURE:
;; OTHER INFORMATION: Description of the artificial sequence: peptide E6 24-38
US-10-476-570-22

Query Match 100.0%; Score 45; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.09; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
|||
Db 3 LQTTIHDI 11

RESULT 9
US-10-476-570-8
; Sequence 8, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 8
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-34
US-10-476-570-8

Query Match 100.0%; Score 45; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.13; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
|||
Db 13 LQTTIHDI 21

RESULT 10
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01553
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980

;; PRIOR FILING DATE: 2001-05-04
;; NUMBER OF SEQ ID NOS: 63
;; SOFTWARE: Patentin Ver. 2.1
;; SEQ ID NO 53
;; LENGTH: 30
;; TYPE: PRT
;; ORGANISM: artificial sequence
;; FEATURE:
;; OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53

Query Match 100.0%; Score 45; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.19; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
|||
Db 12 LQTTIHDI 20

RESULT 11
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match 100.0%; Score 45; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.19; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
|||
Db 12 LQTTIHDI 20

RESULT 12
US-10-476-570-9
; Sequence 9, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570

CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 9
LENGTH: 32
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9

Query Match 100.0%; Score 45; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
DB 13 LQTTIHDI 21

RESULT 13
US-10-476-570-19
Sequence 19, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLAD, Isabelle
APPLICANT: GUILLET, Jean-Gerard
APPLICANT: POUVELLE-MORATILLE, Sandra
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
TITLE OF INVENTION: papillomavirus proteins and uses thereof
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476, 570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 19
LENGTH: 33
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19

Query Match 100.0%; Score 45; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
DB 13 LQTTIHDI 21

RESULT 14
US-10-177-390-6
Sequence 6, Application US/10177390
Publication No. US20030143743A1
GENERAL INFORMATION:
APPLICANT: Schuler, Gerold
APPLICANT: N.V. Antwerp Innovatiecentrum
TITLE OF INVENTION: Improved transfection of Eucaryotic Cells with linear
TITLE OF INVENTION: Polynucleotides by Electroporation
FILE REFERENCE: 021505wo/JH/ml

CURRENT APPLICATION NUMBER: US/10/177,390
CURRENT FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match 100.0%; Score 45; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
DB 19 LQTTIHDI 27

RESULT 15
US-10-484-063-20
Sequence 20, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASSTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILTERMO
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484, 063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306, 809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 20
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-484-063-20

Query Match 100.0%; Score 45; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
DB 19 LQTTIHDI 27

RESULT 16
US-10-484-063-27
Sequence 27, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASSTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILTERMO
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484, 063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306, 809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27

LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match 100.0%; Score 45; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 1.1; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

QY 1 LQTTIHDI 9
DB 19 LQTTIHDI 27

RESULT 17
US-10-858-384-2
Sequence 2, Application US/10858384
Publication No. US20050033025A1
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCINE
APPLICANT: FERRIES, ESTELLE
TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
TITLE OF INVENTION: PARTICULARLY IN VACCINATION
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858,384
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match 100.0%; Score 45; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 1.2; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

QY 1 LQTTIHDI 9
DB 26 LQTTIHDI 34

RESULT 18
US-10-367-057-16
Sequence 16, Application US/10367057
Publication No. US20050100554A1
GENERAL INFORMATION:
APPLICANT: Cuthill, Scott;
APPLICANT: Jackson, Amanda;
APPLICANT: Lewin, David A.;
APPLICANT: Ooi, Chean Eng
TITLE OF INVENTION: Complexes and Methods of Using Same
FILE REFERENCE: 21402-559
CURRENT APPLICATION NUMBER: US/10/367,057
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: 60/256,911
PRIOR FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 198
SOFTWARE: CuraSeqList version 0.1
SEQ ID NO 16
LENGTH: 158
TYPE: PRT
ORGANISM: Homo sapiens
US-10-367-057-16

Query Match 100.0%; Score 45; DB 5; Length 158;

Best Local Similarity 100.0%; Pred. No. 1.2; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

QY 1 LQTTIHDI 9
DB 26 LQTTIHDI 34

RESULT 19
US-11-021-949-13
Sequence 13, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
TITLE OF INVENTION: AND METHODS OF THEIR USE
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 13
LENGTH: 158
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match 100.0%; Score 45; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 1.2; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

QY 1 LQTTIHDI 9
DB 26 LQTTIHDI 34

RESULT 20
US-10-472-724-2
Sequence 2, Application US/10472724
Publication No. US20040171806A1
GENERAL INFORMATION:
APPLICANT: Cid-Arregui, Angel
APPLICANT: Zur Hausen, Harald
TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
FILE REFERENCE: 4121-154
CURRENT APPLICATION NUMBER: US/10/472,724
CURRENT FILING DATE: 2003-09-17
PRIOR APPLICATION NUMBER: PCT/EP02/03271
PRIOR FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: EP 01107271.7
PRIOR FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.2
SEQ ID NO 2
LENGTH: 171
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match 100.0%; Score 45; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 1.3; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

QY 1 LQTTIHDI 9
DB 26 LQTTIHDI 34

Db 31 LQTTIHDI 39

```
RESULT 21
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENV, Marie-Paule
; APPLICANT: BALLOUT, Jean-Marc
; APPLICANT: BILLOUANE, Nadine
; TITLE OF INVENTION: ANTIMICROBIAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; CURRENT FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone B6*TMF.
US-11-072-288-1
```

```
Query Match          100.0%; Score 45; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 1.9;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 LQTTIHDI 9
|||||||
Db 54 LQTTIHDI 62

```
RESULT 22
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Publication No. US20020081329A1
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; CURRENT FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1
```

```
Query Match          100.0%; Score 45; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 LQTTIHDI 9
|||||||
Db 26 LQTTIHDI 34RESULT 23
US-10-000-903-4

```
; Sequence 4, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernarde
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-4
```

```
Query Match          100.0%; Score 45; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 LQTTIHDI 9
|||||||
Db 132 LQTTIHDI 140

```
RESULT 24
US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4
```

```
Query Match          100.0%; Score 45; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 LQTTIHDI 9
|||||||

Db 132 LQTTIHDI 140

RESULT 25
US-10-000-903-10

```
/ Sequence 10, Application US/10000903
/ Publication No. US20020182221A1
/ GENERAL INFORMATION:
/ APPLICANT: Bruck, Claudine
/ APPLICANT: Cabezon Silva, Teresa
/ APPLICANT: Delisse, Anne-Marie Eva Fernande
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ APPLICANT: Lombardo-Bencheikh, Angela
/ TITLE OF INVENTION: Vaccine
/ FILE REFERENCE: B45107
/ CURRENT APPLICATION NUMBER: US/10/000,903
/ CURRENT FILING DATE: 2001-10-01
/ PRIOR APPLICATION NUMBER: PCT/EP98/05285
/ PRIOR FILING DATE: 1998-08-17
/ PRIOR APPLICATION NUMBER: GB 9717953.5
/ PRIOR FILING DATE: 1997-08-22
/ NUMBER OF SEQ ID NOS: 23
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 10
/ LENGTH: 292
/ TYPE: PRT
/ ORGANISM: Homo sapien
US-10-000-903-10
```

```
Query Match 100.0%; Score 45; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 2.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

Qy 1 LQTTIHDI 9
Db 151 LQTTIHDI 159RESULT 26
US-10-899-771-10

```
/ Sequence 10, Application US/10899771
/ Publication No. US20050031638A1
/ GENERAL INFORMATION:
/ APPLICANT: Dalemans, Wilfried L.J.
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
/ TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
/ FILE REFERENCE: B45124
/ CURRENT APPLICATION NUMBER: US/10/899,771
/ CURRENT FILING DATE: 2004-07-27
/ PRIOR APPLICATION NUMBER: US/09/581,976
/ PRIOR FILING DATE: 2000-06-20
/ PRIOR APPLICATION NUMBER: PCT/EP98/08563
/ PRIOR FILING DATE: 1998-12-18
/ PRIOR APPLICATION NUMBER: GB 9727262.9
/ PRIOR FILING DATE: 1997-12-24
/ NUMBER OF SEQ ID NOS: 28
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 10
/ LENGTH: 292
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Chimeric protein (C1ya from Streptococcus
/ OTHER INFORMATION: pneumoniae and B6 from Human papilloma virus type
/ OTHER INFORMATION: 16)
US-10-899-771-10
```

```
Query Match 100.0%; Score 45; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 2.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

Qy 1 LQTTIHDI 9

Db 151 LQTTIHDI 159

RESULT 27
US-10-000-903-6

```
/ Sequence 6, Application US/10000903
/ Publication No. US20020182221A1
/ GENERAL INFORMATION:
/ APPLICANT: Bruck, Claudine
/ APPLICANT: Cabezon Silva, Teresa
/ APPLICANT: Delisse, Anne-Marie Eva Fernande
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ APPLICANT: Lombardo-Bencheikh, Angela
/ TITLE OF INVENTION: Vaccine
/ FILE REFERENCE: B45107
/ CURRENT APPLICATION NUMBER: US/10/000,903
/ CURRENT FILING DATE: 2001-10-01
/ PRIOR APPLICATION NUMBER: PCT/EP98/05285
/ PRIOR FILING DATE: 1998-08-17
/ PRIOR APPLICATION NUMBER: GB 9717953.5
/ PRIOR FILING DATE: 1997-08-22
/ NUMBER OF SEQ ID NOS: 23
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 6
/ LENGTH: 371
/ TYPE: PRT
/ ORGANISM: Homo sapien
US-10-000-903-6
```

```
Query Match 100.0%; Score 45; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

Qy 1 LQTTIHDI 9
Db 132 LQTTIHDI 140RESULT 28
US-10-899-771-6

```
/ Sequence 6, Application US/10899771
/ Publication No. US20050031638A1
/ GENERAL INFORMATION:
/ APPLICANT: Dalemans, Wilfried L.J.
/ APPLICANT: Gerard, Catherine Marie Ghislaine
/ TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
/ TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a CpG Oligonucleotide
/ FILE REFERENCE: B45124
/ CURRENT APPLICATION NUMBER: US/10/899,771
/ CURRENT FILING DATE: 2004-07-27
/ PRIOR APPLICATION NUMBER: US/09/581,976
/ PRIOR FILING DATE: 2000-06-20
/ PRIOR APPLICATION NUMBER: PCT/EP98/08563
/ PRIOR FILING DATE: 1998-12-18
/ PRIOR APPLICATION NUMBER: GB 9727262.9
/ PRIOR FILING DATE: 1997-12-24
/ NUMBER OF SEQ ID NOS: 28
/ SOFTWARE: FastSeq for Windows Version 3.0
/ SEQ ID NO 6
/ LENGTH: 371
/ TYPE: PRT
/ ORGANISM: Artificial Sequence
/ FEATURE:
/ OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
/ OTHER INFORMATION: influenzae B and B67 fusion from Human papilloma
/ OTHER INFORMATION: virus type 16)
US-10-899-771-6
```

```
Query Match 100.0%; Score 45; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 LQTTTHDII 9
| | | | |
Db 132 LQTTTHDII 140

RESULT 29

US-10-000-903-14
; Sequence 14, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fermande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-14

Query Match 100.0%; Score 45; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTTHDII 9
| | | | |
Db 151 LQTTTHDII 159

RESULT 30
US-10-899-771-14
; Sequence 14, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuncted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-14

Query Match 100.0%; Score 45; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 3.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTTHDII 9
| | | | |
Db 151 LQTTTHDII 159

RESULT 31

US-10-367-095-10
; Sequence 10, Application US/10367095
; Publication No. US20030228696A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
; FILE REFERENCE: 44149-1US1
; CURRENT APPLICATION NUMBER: US/10/367,095
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-367-095-10

Query Match 100.0%; Score 45; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 4.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTTHDII 9
| | | | |
Db 496 LQTTTHDII 504

RESULT 32
US-10-368-046-10
; Sequence 10, Application US/10368046
; Publication No. US20040063188A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; APPLICANT: Victoria Cloce
; TITLE OF INVENTION: Method for Isolation and Purification of
; TITLE OF INVENTION: Expressed Gene Products In Vitro
; FILE REFERENCE: 44149-3US1
; CURRENT APPLICATION NUMBER: US/10/368,046
; CURRENT FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118

```

; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-368-046-10

Query Match          100.0%; Score 45; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 4.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 496 LQTTIHDI 504

RESULT 33
; US-10-367-367-10
; Sequence 10, Application US/10367367
; Publication No. US20040121465A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: Optimization of Gene Sequences of
; FILE REFERENCE: 44149-2US1
; CURRENT FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-367-367-10

Query Match          100.0%; Score 45; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 4.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 496 LQTTIHDI 504
```

```

; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-918-337-10

Query Match          100.0%; Score 45; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 4.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 496 LQTTIHDI 504

RESULT 34
; US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:
; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; FILE REFERENCE: 19065/2132
; CURRENT FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
; US-10-918-337-10

Query Match          100.0%; Score 45; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 4.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 496 LQTTIHDI 504

RESULT 35
; US-11-021-949-1
; Sequence 1, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
```

SOFTWARE: FastSeq for windows Version 4.0
; SEQ ID NO 1
; LENGTH: 25
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-1

Query Match 91.1%; Score 41; DB 6; Length 25;
Best Local Similarity 100.0%; Pred. No. 0.91;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
Db 18 LQTTIHDI 25

RESULT 36
US-10-820-155-42
; Sequence 42, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: Nactimmune A/S
; APPLICANT: Wellguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; CURRENT FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: Patent in version 3.1
; SEQ ID NO 42
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Canis familiaris
US-10-820-155-42

Query Match 82.2%; Score 37; DB 5; Length 248;
Best Local Similarity 75.0%; Pred. No. 64;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
Db 107 LQTTIHDI 114

RESULT 37
US-10-820-155-57
; Sequence 57, Application US/10820155
; Publication No. US20050137126A1
; GENERAL INFORMATION:
; APPLICANT: Nactimmune A/S
; APPLICANT: Wellguny, Dietmar
; APPLICANT: Jensenius, Jens Christian
; APPLICANT: Kongerslev, Leif
; APPLICANT: Mathiesen, Finn
; TITLE OF INVENTION: Treatment of SARS in individuals
; FILE REFERENCE: P 774 US00
; CURRENT APPLICATION NUMBER: US/10/820,155
; CURRENT FILING DATE: 2004-04-08
; NUMBER OF SEQ ID NOS: 137
; SOFTWARE: Patent in version 3.1
; SEQ ID NO 57
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Canis familiaris
US-10-820-155-57

Query Match 82.2%; Score 37; DB 5; Length 248;
Best Local Similarity 75.0%; Pred. No. 64;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
Db 107 LQTTIHDI 114

RESULT 38
US-10-476-570-23
; Sequence 23, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05380
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: Patent in Ver. 2.1
; SEQ ID NO 23
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 28-42
US-10-476-570-23

Query Match 80.0%; Score 36; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 4.7;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 3 TTTHDII 9
Db 1 TTTHDII 7

RESULT 39
US-10-425-115-237192
; Sequence 237192, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 237192
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(151)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_147905C.1.pep
US-10-425-115-237192

Query Match 80.0%; Score 36; DB 4; Length 151;
Best Local Similarity 75.0%; Pred. No. 58;

Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 8
|:|:|:|
Db 96 LTTIHDI 103

RESULT 40

US-11-097-143-4221
; Sequence 4221, Application US/11097143
; Publication No. US200502085581
; GENERAL INFORMATION:
; APPLICANT: Venter, J. Craig
; APPLICANT: et al.
; TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
; TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
; TITLE OF INVENTION: DROSOPHILA GENES.
; FILE REFERENCE: CLO00728
; CURRENT APPLICATION NUMBER: US/11/097,143
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: 60/157,832
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: 60/160,191
; PRIOR FILING DATE: 1999-10-19
; PRIOR APPLICATION NUMBER: 60/161,932
; PRIOR FILING DATE: 1999-10-28
; PRIOR APPLICATION NUMBER: 60/164,769
; PRIOR FILING DATE: 1999-11-12
; PRIOR APPLICATION NUMBER: 60/173,383
; PRIOR FILING DATE: 1999-12-28
; PRIOR APPLICATION NUMBER: 60/175,693
; PRIOR FILING DATE: 2000-01-12
; PRIOR APPLICATION NUMBER: 60/184,831
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60/191,637
; PRIOR FILING DATE: 2000-03-23
; NUMBER OF SEQ ID NOS: 43008
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 4221
; LENGTH: 601
; TYPE: PRT
; ORGANISM: DROSOPHILA
US-11-097-143-4221

Query Match 80.0%; Score 36; DB 6; Length 601;

Best Local Similarity 100.0%; Pred. No. 2.6e+02;

Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 3 TTTIHDI 9
|:|:|:|
Db 273 TTTIHDI 279

RESULT 41

US-10-282-122A-49664
; Sequence 49664, Application US/10282122A
; Publication No. US20040029129A1
; GENERAL INFORMATION:
; APPLICANT: Wang, Lianguu
; APPLICANT: Zamudio, Carlos
; APPLICANT: Malone, Cheryl
; APPLICANT: Haseelbeck, Robert
; APPLICANT: Ohlsen, Karl
; APPLICANT: Zyskind, Judith
; APPLICANT: Wall, Daniel
; APPLICANT: Trawick, John
; APPLICANT: Carr, Grant
; APPLICANT: Yamamoto, Robert
; APPLICANT: Foreyth, R.
; APPLICANT: Xu, H.
; TITLE OF INVENTION: Identification of Essential Genes in Microorganisms
; FILE REFERENCE: EUTTRA.034A
; CURRENT APPLICATION NUMBER: US/10/282,122A

CURRENT FILING DATE: 2003-02-20

PRIOR APPLICATION NUMBER: 60/191,078

PRIOR FILING DATE: 2000-03-21

PRIOR APPLICATION NUMBER: 60/206,848

PRIOR FILING DATE: 2000-05-23

PRIOR APPLICATION NUMBER: 60/207,727

PRIOR FILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: 60/230,335

PRIOR FILING DATE: 2000-09-06

PRIOR APPLICATION NUMBER: 60/230,347

PRIOR FILING DATE: 2000-09-09

PRIOR APPLICATION NUMBER: 60/242,578

PRIOR FILING DATE: 2000-10-23

PRIOR APPLICATION NUMBER: 60/253,625

PRIOR FILING DATE: 2000-11-27

PRIOR APPLICATION NUMBER: 60/257,931

PRIOR FILING DATE: 2000-12-22

PRIOR APPLICATION NUMBER: 60/267,636

PRIOR FILING DATE: 2001-02-09

PRIOR APPLICATION NUMBER: 60/269,308

PRIOR FILING DATE: 2001-02-16

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 78614

SOFTWARE: PatentIn version 3.1

SEQ ID NO 49664

LENGTH: 1082

TYPE: PRT

ORGANISM: Burkholderia fungorum

US-10-282-122A-49664

Query Match 80.0%; Score 36; DB 4; Length 1082;

Best Local Similarity 55.6%; Pred. No. 5e+02;

Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
|:|:|:|
Db 620 LDTVHDVL 628

RESULT 42

US-10-452-024-95
; Sequence 95, Application US/10452024
; Publication No. US20040013687A1
; GENERAL INFORMATION:
; APPLICANT: Simpson, Lance
; APPLICANT: Park, Jung-Beak
; APPLICANT: Maksymowich, Andrew
; TITLE OF INVENTION: Compositions and Methods for Trans epithelial Molecular Transport
; FILE REFERENCE: 9855-9601
; CURRENT APPLICATION NUMBER: US/10/452,024
; CURRENT FILING DATE: 2003-06-02
; PRIOR APPLICATION NUMBER: 60/384,949
; PRIOR FILING DATE: 2002-05-31
; NUMBER OF SEQ ID NOS: 188
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 95
; LENGTH: 1198
; TYPE: PRT
; ORGANISM: Clostridium botulinum
US-10-452-024-95

Query Match 80.0%; Score 36; DB 4; Length 1198;

Best Local Similarity 85.7%; Pred. No. 5.5e+02;

Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 2 QTTIHDI 8
|:|:|:|
Db 325 QTTVHDI 331

RESULT 43

US-11-097-143-36795
; Sequence 36795, Application US/11097143

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; Publication No. US20050208558A1
; GENERAL INFORMATION:
; APPLICANT: Venter, J. Craig
; TITLE OF INVENTION: DETECTION KIT, SUCH AS NUCLEIC ACID
; TITLE OF INVENTION: ARRAYS, FOR DETECTING EXPRESSION OF 10,000 OR MORE
; FILE REFERENCE: CLO00728
; CURRENT APPLICATION NUMBER: US/11/097,143
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: 60/157,832
; PRIOR FILING DATE: 1999-10-05
; PRIOR APPLICATION NUMBER: 60/160,191
; PRIOR FILING DATE: 1999-10-19
; PRIOR APPLICATION NUMBER: 60/161,932
; PRIOR FILING DATE: 1999-10-28
; PRIOR APPLICATION NUMBER: 60/164,769
; PRIOR FILING DATE: 1999-11-12
; PRIOR APPLICATION NUMBER: 60/173,383
; PRIOR FILING DATE: 1999-12-28
; PRIOR APPLICATION NUMBER: 60/175,693
; PRIOR FILING DATE: 2000-01-12
; PRIOR APPLICATION NUMBER: 60/184,831
; PRIOR FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: 60/191,637
; PRIOR FILING DATE: 2000-03-23
; NUMBER OF SEQ ID NOS: 43008
; SOFTWARE: PASTESEQ for Windows Version 4.0
; SEQ ID NO 36795
; LENGTH: 1381
; TYPE: PRT
; ORGANISM: DROSOPHILA
; US-11-097-143-36795
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Query Match      80.0%; Score 36; DB 6; Length 1381;
Best Local Similarity 66.7%; Pred. No. 6; Se+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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QY 1 LQTTIHDI 9
DB 366 LSTIHDLI 374
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RESULT 44
US-10-767-701-50864
; Sequence 50864, Application US/10767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(5335)B
; CURRENT APPLICATION NUMBER: US/10/767,701
; CURRENT FILING DATE: 2004-01-29
; NUMBER OF SEQ ID NOS: 63128
; SEQ ID NO 50864
; LENGTH: 97
; TYPE: PRT
; ORGANISM: Sorghum bicolor
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB3480-038-P1-K1-D12.pep
; US-10-767-701-50864
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Query Match      77.8%; Score 35; DB 4; Length 97;
Best Local Similarity 66.7%; Pred. No. 55;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
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QY 1 LQTTIHDI 9
DB 8 VKTTHDIM 16
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RESULT 45
US-10-425-115-263392
; Sequence 263392, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53322)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 263392
; LENGTH: 156
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(156)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MFT4577_171828C.1.pep
; US-10-425-115-263392
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Query Match      77.8%; Score 35; DB 4; Length 156;
Best Local Similarity 66.7%; Pred. No. 93;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
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QY 1 LQTTIHDI 9
DB 67 VKTTHDIM 75
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RESULT 46
US-10-425-114-55470
; Sequence 55470, Application US/10425114
; Publication No. US2004003488A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jindong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven B
; APPLICANT: Tabaska, Jack B
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 55470
; LENGTH: 165
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-GMFLMINISOY117C04_FLI.pep
; US-10-425-114-55470
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Query Match      77.8%; Score 35; DB 4; Length 165;
Best Local Similarity 66.7%; Pred. No. 99;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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QY 1 LQTTIHDI 9
DB 125 LQTSIHDFI 133
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RESULT 47
US-10-424-599-168462
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; Sequence 168462, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 168462
; LENGTH: 316
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(316)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_123136C.1.pep
US-10-424-599-168462

Query Match          77.8%; Score 35; DB 4; Length 316;
Best Local Similarity 66.7%; Pred. No. 2e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY      1 LQTTIHDI 9
Db      123 INTIHDI 131

RESULT 48
US-10-424-599-230648
; Sequence 230648, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 230648
; LENGTH: 336
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_50297C.1.pep
US-10-424-599-230648

Query Match          77.8%; Score 35; DB 4; Length 336;
Best Local Similarity 66.7%; Pred. No. 2.1e+02;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY      1 LQTTIHDI 9
Db      210 LQTSIHDI 218

RESULT 49
US-10-437-963-117225
; Sequence 117225, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
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; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 117225
; LENGTH: 403
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_20650C.1.pep
US-10-437-963-117225

Query Match          77.8%; Score 35; DB 4; Length 403;
Best Local Similarity 66.7%; Pred. No. 2.6e+02;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

OY      1 LQTTIHDI 9
Db      234 VKTTIHDI 242

RESULT 50
US-10-425-115-263390
; Sequence 263390, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 263390
; LENGTH: 448
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_171826C.1.pep
US-10-425-115-263390

Query Match          77.8%; Score 35; DB 4; Length 448;
Best Local Similarity 66.7%; Pred. No. 2.9e+02;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

OY      1 LQTTIHDI 9
Db      255 VKTTIHDI 263

Search completed: May 5, 2006, 08:07:01
Job time : 65 secs
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GenCore version 5.1.7
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OM protein - protein search, using sw model1

Run on: May 5, 2006, 07:56:56 ; Search time 8.4 Seconds
(Without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-6
Perfect score: 45
Sequence: 1 LGTHIDII 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

Published Applications AA New:*
1: /SIDSS/ptodata/1/pubppaa/US08_NEW_PUB.pep1:*
2: /SIDSS/ptodata/1/pubppaa/US06_NEW_PUB.pep1:*
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12: /SIDSS/ptodata/1/pubppaa/US60_NEW_PUB.pep1:*

Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	45	100.0	151	9	US-10-530-253-13
2	45	100.0	158	11	US-11-206-138-3
3	45	100.0	248	9	US-10-530-253-1
4	45	100.0	248	9	US-10-530-253-3
5	45	100.0	248	9	US-10-530-253-5
6	45	100.0	248	9	US-10-530-253-7
7	45	100.0	248	9	US-10-530-253-9
8	45	100.0	248	9	US-10-530-253-11
9	45	100.0	256	11	US-11-192-923A-2
10	35	77.8	250	11	US-11-096-568A-1943
11	35	77.8	287	11	US-11-096-568A-19577
12	35	77.8	298	11	US-11-096-568A-1942
13	35	77.8	303	11	US-11-096-568A-1941
14	35	77.8	329	11	US-11-096-568A-1940
15	35	77.8	448	11	US-11-096-568A-19576
16	35	77.8	463	11	US-11-096-568A-19575
17	35	77.8	540	11	US-11-096-568A-19579
18	35	77.8	989	11	US-11-188-298-5563
19	33	73.3	149	9	US-10-530-253-17
20	33	73.3	158	9	US-10-530-253-26
21	33	73.3	237	9	US-10-506-454-1603

22	71.1	333	11	US-11-188-298-17532	Sequence 17532, A
23	71.1	337	11	US-11-188-298-16137	Sequence 16137, A
24	71.1	340	11	US-11-188-298-6731	Sequence 6731, Ap
25	71.1	513	9	US-10-873-528-37	Sequence 37, Appl
26	71.1	513	9	US-10-873-528-193	Sequence 193, Appl
27	71.1	553	11	US-11-103-957-61	Sequence 61, Appl
28	71.1	643	11	US-11-087-099-1991	Sequence 1991, Ap
29	68.9	186	11	US-11-087-099-10391	Sequence 10391, A
30	68.9	187	11	US-11-087-099-7508	Sequence 7508, Ap
31	68.9	335	11	US-11-188-298-21765	Sequence 21765, A
32	68.9	340	11	US-11-188-298-16409	Sequence 16409, A
33	68.9	333	9	US-10-703-799B-62	Sequence 62, Appl
34	66.7	19	9	US-10-503-575-174	Sequence 174, Appl
35	66.7	63	11	US-11-079-463-6103	Sequence 6103, Ap
36	66.7	133	9	US-10-986-405-199	Sequence 199, App
37	66.7	135	9	US-10-986-405-205	Sequence 205, App
38	66.7	140	9	US-10-986-405-229	Sequence 229, App
39	66.7	149	9	US-10-986-405-200	Sequence 200, App
40	66.7	179	11	US-11-188-298-10295	Sequence 10295, A
41	66.7	184	9	US-10-986-405-216	Sequence 216, App
42	66.7	221	11	US-11-188-298-17244	Sequence 17244, A
43	66.7	333	11	US-11-188-298-3948	Sequence 3948, Ap
44	66.7	333	11	US-11-188-298-22053	Sequence 22053, A
45	66.7	334	11	US-11-188-298-2051	Sequence 2051, Ap
46	66.7	338	11	US-11-096-568A-21865	Sequence 21865, A
47	66.7	352	11	US-11-188-298-12429	Sequence 12429, A
48	66.7	354	9	US-10-878-556A-84	Sequence 84, Appl
49	66.7	354	11	US-11-100-640-4	Sequence 4, Appl1
50	66.7	371	11	US-11-079-463-6883	Sequence 6883, Ap
51	66.7	419	11	US-11-096-568A-21864	Sequence 21864, A
52	66.7	470	11	US-11-079-463-6501	Sequence 6501, Ap
53	66.7	474	11	US-11-096-568A-21863	Sequence 21863, A
54	66.7	522	11	US-11-087-099-2820	Sequence 2820, Ap
55	66.7	522	11	US-11-188-298-2685	Sequence 2685, Ap
56	66.7	550	11	US-11-072-512-3416	Sequence 3416, Ap
57	66.7	725	11	US-11-078-189-15	Sequence 15, Appl
58	66.7	868	9	US-10-821-234-1082	Sequence 1082, Ap
59	66.7	1453	11	US-11-096-568A-14692	Sequence 14692, A
60	66.7	1473	11	US-11-096-568A-14691	Sequence 14691, A
61	66.7	1529	11	US-11-096-568A-14690	Sequence 14690, A
62	66.7	1763	8	US-10-504-120-21	Sequence 21, Appl
63	66.7	1807	8	US-10-504-120-22	Sequence 22, Appl
64	66.7	15	9	US-10-530-061-1673	Sequence 1673, Ap
65	64.4	37	9	US-10-729-121-22	Sequence 22, Appl
66	64.4	37	11	US-11-285-537-22	Sequence 22, Appl
67	64.4	149	9	US-10-530-253-24	Sequence 24, Appl
68	64.4	165	9	US-10-793-626-2528	Sequence 2528, Ap
69	64.4	169	11	US-11-183-664-12	Sequence 12, Appl
70	64.4	194	11	US-11-188-298-2828	Sequence 2828, Ap
71	64.4	195	11	US-11-188-298-578	Sequence 578, App
72	64.4	203	11	US-11-188-298-15244	Sequence 15244, A
73	64.4	207	11	US-11-188-298-20708	Sequence 20708, A
74	64.4	215	11	US-11-183-664-26	Sequence 26, Appl
75	64.4	218	11	US-11-096-568A-15664	Sequence 15664, A
76	64.4	223	11	US-11-188-298-23212	Sequence 23212, A
77	64.4	289	11	US-11-045-004-993	Sequence 993, App
78	64.4	295	11	US-11-188-298-12304	Sequence 12304, A
79	64.4	295	11	US-11-188-298-20727	Sequence 20727, A
80	64.4	315	11	US-11-188-298-11316	Sequence 11316, A
81	64.4	317	11	US-11-188-298-17553	Sequence 17553, A
82	64.4	329	11	US-11-188-298-19388	Sequence 19388, A
83	64.4	330	11	US-11-188-298-9402	Sequence 9402, Ap
84	64.4	331	11	US-11-188-298-12669	Sequence 12669, A
85	64.4	331	11	US-11-188-298-15348	Sequence 15348, A
86	64.4	331	11	US-11-188-298-15995	Sequence 15995, A
87	64.4	331	11	US-11-188-298-18595	Sequence 18595, A
88	64.4	333	11	US-11-188-298-18079	Sequence 18079, A
89	64.4	334	11	US-11-188-298-15372	Sequence 15372, A
90	64.4	335	11	US-11-188-298-1265	Sequence 1265, Ap
91	64.4	335	11	US-11-188-298-9411	Sequence 9411, Ap
92	64.4	336	11	US-11-188-298-20606	Sequence 20606, A
93	64.4	336	11	US-11-188-298-22529	Sequence 22529, A
94	64.4	337	11	US-11-188-298-9863	Sequence 9863, A

Tue May 9 09:28:57 2006

us-08-170-344-6.rapbn

95	29	64.4	337	11	US-11-188-298-13243	Sequence 13243, A
96	29	64.4	337	11	US-11-188-298-15095	Sequence 15095, A
97	29	64.4	337	11	US-11-188-298-17839	Sequence 17839, A
98	29	64.4	338	11	US-11-188-298-1497	Sequence 1497, Ap
99	29	64.4	338	11	US-11-188-298-8080	Sequence 8080, Ap
100	29	64.4	338	11	US-11-188-298-14046	Sequence 14046, A
101	29	64.4	338	11	US-11-188-298-14254	Sequence 14254, A
102	29	64.4	338	11	US-11-188-298-15011	Sequence 15011, A
103	29	64.4	338	11	US-11-188-298-15408	Sequence 15408, A
104	29	64.4	338	11	US-11-188-298-18267	Sequence 18267, A
105	29	64.4	338	11	US-11-188-298-18443	Sequence 18443, A
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107	29	64.4	340	11	US-11-188-298-1105	Sequence 1105, Ap
108	29	64.4	340	11	US-11-188-298-1507	Sequence 1507, Ap
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110	29	64.4	340	11	US-11-188-298-6179	Sequence 6179, Ap
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112	29	64.4	341	11	US-11-188-298-6229	Sequence 6229, A
113	29	64.4	341	11	US-11-188-298-6225	Sequence 6225, A
114	29	64.4	343	11	US-11-188-298-17937	Sequence 17937, A
115	29	64.4	345	11	US-11-188-298-17029	Sequence 20, Ap1
116	29	64.4	348	11	US-11-188-298-11222	Sequence 17029, A
117	29	64.4	351	11	US-11-188-298-2278	Sequence 2278, A
118	29	64.4	357	11	US-11-096-568A-23149	Sequence 2278, Ap
119	29	64.4	357	11	US-11-188-298-1920	Sequence 1920, A
120	29	64.4	357	11	US-11-188-298-1920	Sequence 1920, A
121	29	64.4	360	11	US-11-188-298-10689	Sequence 10689, A
122	29	64.4	364	11	US-11-096-568A-23148	Sequence 23148, A
123	29	64.4	380	9	US-10-530-773-443	Sequence 443, Ap
124	29	64.4	402	11	US-11-096-568A-23147	Sequence 23147, A
125	29	64.4	437	9	US-11-188-298-14579	Sequence 14579, A
126	29	64.4	437	9	US-10-821-234-1392	Sequence 1392, Ap
127	29	64.4	437	9	US-11-188-298-21184	Sequence 21184, A
128	29	64.4	437	9	US-10-770-726-72	Sequence 72, Ap1
129	29	64.4	437	9	US-11-096-568A-34055	Sequence 34055, A
130	29	64.4	437	9	US-11-031-206-106	Sequence 106, Ap
131	29	64.4	437	9	US-11-096-568A-34054	Sequence 34054, A
132	29	64.4	437	9	US-11-096-568A-34053	Sequence 34053, A
133	29	64.4	437	9	US-11-080-991-108	Sequence 108, Ap
134	29	64.4	437	9	US-11-188-298-18205	Sequence 18205, A
135	29	64.4	437	9	US-11-188-298-301-20	Sequence 20, Ap1
136	29	64.4	437	9	US-11-188-298-301-19	Sequence 19, Ap1
137	29	64.4	437	9	US-11-188-298-301-21	Sequence 21, Ap1
138	29	64.4	437	9	US-11-188-298-3797	Sequence 3797, Ap
139	29	64.4	437	9	US-11-045-004-266	Sequence 266, Ap
140	29	64.4	437	9	US-11-189-301-10	Sequence 10, Ap1
141	29	64.4	437	9	US-10-915-002-321	Sequence 321, Ap
142	29	64.4	437	9	US-10-915-002-322	Sequence 322, Ap
143	29	64.4	437	9	US-10-915-002-323	Sequence 323, Ap
144	29	64.4	437	9	US-10-204-639-4	Sequence 4, Ap11
145	29	64.4	437	9	US-10-204-639-54	Sequence 54, Ap1
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147	29	64.4	437	9	US-10-877-346-44	Sequence 44, Ap1
148	29	64.4	437	9	US-10-995-561-593	Sequence 593, Ap
149	29	64.4	437	9	US-10-816-768-46	Sequence 46, Ap1
150	29	64.4	437	9	US-11-226-555-12	Sequence 12, Ap1
151	29	64.4	437	9	US-11-096-568A-20906	Sequence 20906, A
152	29	64.4	437	9	US-11-096-568A-20905	Sequence 20905, A
153	29	64.4	437	9	US-11-188-298-9723	Sequence 9723, Ap
154	29	64.4	437	9	US-11-087-099-1865	Sequence 1865, Ap
155	29	64.4	437	9	US-11-087-099-2617	Sequence 2617, Ap
156	29	64.4	437	9	US-11-045-004-983	Sequence 983, Ap
157	29	64.4	437	9	US-11-045-004-375	Sequence 375, Ap
158	29	64.4	437	9	US-11-096-568A-20904	Sequence 20904, A
159	29	64.4	437	9	US-10-467-657-2290	Sequence 2290, Ap
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161	29	64.4	437	9	US-10-194-487-374	Sequence 374, Ap
162	29	64.4	437	9	US-10-195-883-374	Sequence 374, Ap
163	29	64.4	437	9	US-10-195-888-374	Sequence 374, Ap
164	29	64.4	437	9	US-10-195-889-374	Sequence 374, Ap
165	29	64.4	437	9	US-10-506-454-1219	Sequence 1219, Ap
166	29	64.4	437	9	US-11-096-686-10545	Sequence 10545, A
167	29	64.4	437	9	US-11-096-686-10545	Sequence 10545, A

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243	27	60.0	331	11	US-11-188-298-3201	Sequence 3201, Ap	316	27	60.0	342	11	US-11-188-298-10269	Sequence 10269, A
244	27	60.0	331	11	US-11-188-298-3228	Sequence 3228, Ap	317	27	60.0	342	11	US-11-188-298-15139	Sequence 15139, A
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247	27	60.0	332	11	US-11-188-298-13876	Sequence 13876, A	320	27	60.0	345	11	US-11-188-298-16881	Sequence 16881, A
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252	27	60.0	335	8	US-10-511-937-2560	Sequence 2560, Ap	325	27	60.0	350	11	US-11-096-568A-1986	Sequence 31986, A
253	27	60.0	335	9	US-10-995-561-704	Sequence 704, App	326	27	60.0	350	11	US-11-188-298-13989	Sequence 12989, A
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258	27	60.0	337	11	US-11-188-298-1785	Sequence 1785, App	331	27	60.0	354	11	US-11-096-568A-6417	Sequence 6417, Ap
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266	27	60.0	337	11	US-11-188-298-5226	Sequence 5226, Ap	339	27	60.0	385	11	US-11-188-298-10855	Sequence 10855, A
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268	27	60.0	337	11	US-11-188-298-7167	Sequence 7167, Ap	341	27	60.0	386	11	US-11-188-298-20493	Sequence 20493, A
269	27	60.0	337	11	US-11-188-298-7573	Sequence 7573, Ap	342	27	60.0	389	11	US-11-096-568A-6415	Sequence 6415, Ap
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273	27	60.0	337	11	US-11-188-298-8881	Sequence 8881, Ap	346	27	60.0	400	11	US-11-096-568A-9720	Sequence 9720, Ap
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295	27	60.0	338	11	US-11-096-568A-9722	Sequence 9722, Ap	368	27	60.0	489	11	US-11-188-298-16624	Sequence 16624, A
296	27	60.0	338	11	US-11-188-298-3869	Sequence 3869, Ap	369	27	60.0	491	11	US-11-188-298-13154	Sequence 13154, A
297	27	60.0	338	11	US-11-188-298-5280	Sequence 5280, Ap	370	27	60.0	496	11	US-11-188-298-9939	Sequence 9939, Ap
298	27	60.0	338	11	US-11-188-298-8233	Sequence 8233, Ap	371	27	60.0	498	11	US-11-188-298-3477	Sequence 3477, Ap
299	27	60.0	338	11	US-11-188-298-10456	Sequence 10456, A	372	27	60.0	514	8	US-10-505-928-335	Sequence 335, App
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311	27	60.0	341	9	US-10-793-626-1188	Sequence 1188, Ap	384	27	60.0	682	11	US-11-045-004-1951	Sequence 1951, Ap
312	27	60.0	341	11	US-11-188-298-8705	Sequence 8705, Ap	385	27	60.0	713	8	US-10-505-928-345	Sequence 345, App
313	27	60.0	341	11	US-11-188-298-8949	Sequence 8949, Ap	386	27	60.0	720	8	US-10-505-928-288	Sequence 288, App

387	27	60.0	819	11	US-11-045-004-778	Sequence 778, App	460	26	57.8	260	11	US-11-188-298-6417	Sequence 6417, App
388	27	60.0	820	11	US-11-147-047-31	Sequence 31, App1	461	26	57.8	261	11	US-11-188-298-6910	Sequence 6910, App
389	27	60.0	860	11	US-11-019-711-59	Sequence 59, App1	462	26	57.8	262	11	US-11-188-298-2240	Sequence 2240, App
390	27	60.0	931	11	US-11-019-711-117	Sequence 117, App	463	26	57.8	263	11	US-11-188-298-9786	Sequence 9786, App
391	27	60.0	931	11	US-11-019-711-118	Sequence 118, App	464	26	57.8	267	11	US-11-188-298-15650	Sequence 15650, App
392	27	60.0	931	11	US-11-019-711-119	Sequence 119, App	465	26	57.8	269	9	US-10-506-454-218	Sequence 218, App
393	27	60.0	931	11	US-11-019-711-120	Sequence 120, App	466	26	57.8	270	11	US-11-188-298-22192	Sequence 22192, App
394	27	60.0	931	11	US-11-183-136-22	Sequence 22, App1	467	26	57.8	271	11	US-11-096-568A-8711	Sequence 8711, App
395	27	60.0	931	11	US-11-183-136-24	Sequence 24, App1	468	26	57.8	271	11	US-11-188-298-11597	Sequence 11697, App
396	27	60.0	967	8	US-10-505-928-795	Sequence 795, App	469	26	57.8	271	11	US-11-045-004-363	Sequence 363, App
397	27	60.0	967	11	US-11-054-281-74	Sequence 74, App1	470	26	57.8	273	11	US-11-188-298-1754	Sequence 12877, App
398	27	60.0	1123	11	US-11-096-568A-27581	Sequence 27581, A	471	26	57.8	273	11	US-11-188-298-12817	Sequence 12817, App
399	27	60.0	1236	11	US-11-199-544-69	Sequence 69, App1	472	26	57.8	274	11	US-11-096-568A-25841	Sequence 25841, App
400	27	60.0	1236	11	US-11-096-568A-27580	Sequence 27580, A	473	26	57.8	275	11	US-11-188-298-9173	Sequence 9173, App
401	27	60.0	1240	11	US-11-091-643-18	Sequence 18, App1	474	26	57.8	276	11	US-11-096-568A-16122	Sequence 16122, A
402	27	60.0	1332	10	US-11-314-018-18	Sequence 18, App1	475	26	57.8	276	11	US-11-096-568A-16633	Sequence 25617, A
403	27	60.0	1332	10	US-11-091-643-18	Sequence 20, App1	476	26	57.8	276	11	US-11-096-568A-25617	Sequence 17192, A
404	27	60.0	1344	10	US-11-314-018-20	Sequence 20, App1	477	26	57.8	276	11	US-11-188-298-14055	Sequence 14055, A
405	27	60.0	1344	10	US-11-091-643-20	Sequence 6, App1	478	26	57.8	276	11	US-11-188-298-14055	Sequence 16201, A
406	27	60.0	1386	10	US-11-314-018-6	Sequence 6, App1	479	26	57.8	277	11	US-11-188-298-16201	Sequence 5324, App
407	27	60.0	1386	11	US-11-091-643-6	Sequence 48, App1	480	26	57.8	278	11	US-11-188-298-8452	Sequence 8452, App
408	27	60.0	1437	11	US-11-014-147-48	Sequence 48, App1	481	26	57.8	278	11	US-11-188-298-9379	Sequence 9379, App
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410	27	60.0	1766	9	US-10-204-639-36	Sequence 36, App1	483	26	57.8	278	11	US-11-188-298-11471	Sequence 11471, A
411	27	60.0	4059	8	US-10-501-834-6	Sequence 6, App1	484	26	57.8	278	11	US-11-188-298-12285	Sequence 12285, A
412	27	60.0	4074	8	US-10-501-834-2	Sequence 2, App1	485	26	57.8	278	11	US-11-188-298-12285	Sequence 13734, A
413	26	57.8	18	11	US-11-145-861-425	Sequence 425, App	486	26	57.8	278	11	US-11-188-298-13734	Sequence 14545, A
414	26	57.8	56	11	US-11-264-096-972	Sequence 972, App	487	26	57.8	278	11	US-11-188-298-14545	Sequence 15010, A
415	26	57.8	71	11	US-11-264-096-1508	Sequence 1508, App	488	26	57.8	278	11	US-11-188-298-15010	Sequence 18326, A
416	26	57.8	75	11	US-11-188-298-14911	Sequence 14911, A	489	26	57.8	278	11	US-11-188-298-19342	Sequence 19342, A
417	26	57.8	78	11	US-11-079-463-6620	Sequence 6620, App	490	26	57.8	278	11	US-11-188-298-20428	Sequence 20428, A
418	26	57.8	114	11	US-11-229-769-321	Sequence 67, App1	491	26	57.8	278	11	US-11-188-298-21172	Sequence 21172, A
419	26	57.8	118	11	US-11-098-686-67	Sequence 67, App1	492	26	57.8	278	11	US-11-188-298-21796	Sequence 21796, A
420	26	57.8	119	11	US-11-087-099-8505	Sequence 8505, App	493	26	57.8	278	11	US-10-506-454-1033	Sequence 1033, App
421	26	57.8	139	11	US-11-188-298-15597	Sequence 15597, A	494	26	57.8	279	9	US-11-079-463-8374	Sequence 8374, App
422	26	57.8	142	9	US-10-467-657-1558	Sequence 1558, App	495	26	57.8	280	11	US-11-188-298-6745	Sequence 6745, App
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426	26	57.8	156	9	US-10-530-253-19	Sequence 19, App1	499	26	57.8	280	11	US-11-188-298-8158	Sequence 82143, App
427	26	57.8	161	11	US-11-188-298-5974	Sequence 5974, App	500	26	57.8	282	11	US-11-096-568A-25840	Sequence 25840, A
428	26	57.8	161	11	US-11-045-004-455	Sequence 455, App	501	26	57.8	282	11	US-11-096-568A-26105	Sequence 26105, A
429	26	57.8	170	11	US-11-079-463-9959	Sequence 9959, App	502	26	57.8	283	11	US-11-096-568A-16632	Sequence 16632, App
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431	26	57.8	176	11	US-11-079-463-6744	Sequence 6744, App	504	26	57.8	299	11	US-11-096-568A-25616	Sequence 25616, A
432	26	57.8	180	11	US-11-232-406A-2	Sequence 2, App1	505	26	57.8	300	11	US-11-188-298-6041	Sequence 6041, App
433	26	57.8	181	9	US-10-485-517-338	Sequence 338, App	506	26	57.8	300	11	US-11-188-298-6041	Sequence 11208, App
434	26	57.8	181	11	US-11-188-298-11483	Sequence 11483, A	507	26	57.8	300	11	US-11-188-298-11208	Sequence 119834, A
435	26	57.8	182	9	US-10-485-517-328	Sequence 328, App	508	26	57.8	300	11	US-11-188-298-11208	Sequence 16, App1
436	26	57.8	186	11	US-11-087-099-12094	Sequence 12094, A	509	26	57.8	301	11	US-11-188-298-11208	Sequence 2591, App
437	26	57.8	190	11	US-11-096-568A-16124	Sequence 16124, A	510	26	57.8	301	11	US-11-134-795-16	Sequence 7458, App
438	26	57.8	190	11	US-11-096-568A-26107	Sequence 26107, A	511	26	57.8	302	11	US-11-188-298-7458	Sequence 15725, A
439	26	57.8	193	11	US-11-087-099-9444	Sequence 9444, App	512	26	57.8	302	11	US-11-188-298-7458	Sequence 16815, A
440	26	57.8	202	11	US-11-072-512-2736	Sequence 2736, App	513	26	57.8	305	11	US-11-188-298-16915	Sequence 21349, A
441	26	57.8	203	11	US-11-087-099-8547	Sequence 8547, App	514	26	57.8	305	11	US-11-188-298-16915	Sequence 5553, App
442	26	57.8	213	11	US-11-079-463-8765	Sequence 8765, App	515	26	57.8	309	11	US-11-188-298-12572	Sequence 12572, App
443	26	57.8	220	11	US-11-096-568A-19911	Sequence 19911, A	516	26	57.8	309	11	US-11-188-298-14834	Sequence 14834, A
444	26	57.8	220	11	US-11-188-298-19130	Sequence 19130, A	517	26	57.8	309	11	US-11-188-298-14834	Sequence 19091, App
445	26	57.8	228	11	US-11-055-822-236	Sequence 236, App	518	26	57.8	309	11	US-11-188-298-14834	Sequence 5845, App
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447	26	57.8	228	11	US-11-055-822-766	Sequence 766, App	520	26	57.8	310	11	US-11-188-298-14005	Sequence 14005, A
448	26	57.8	231	11	US-11-239-674-114	Sequence 114, App	521	26	57.8	310	11	US-11-188-298-14005	Sequence 15179, A
449	26	57.8	231	11	US-11-096-568A-16123	Sequence 16123, A	522	26	57.8	310	11	US-11-188-298-14005	Sequence 19139, A
450	26	57.8	231	11	US-11-096-568A-25618	Sequence 25618, A	523	26	57.8	310	11	US-11-188-298-14005	Sequence 21225, A
451	26	57.8	231	11	US-11-096-568A-25842	Sequence 25842, A	524	26	57.8	310	11	US-11-188-298-14005	Sequence 22360, A
452	26	57.8	231	11	US-11-096-568A-26106	Sequence 26106, A	525	26	57.8	310	11	US-11-188-298-14005	Sequence 13049, A
453	26	57.8	231	11	US-11-045-004-1371	Sequence 1371, App	526	26	57.8	311	11	US-11-188-298-13308	Sequence 13308, A
454	26	57.8	242	11	US-11-087-099-8891	Sequence 8891, App	527	26	57.8	311	11	US-11-188-298-13308	Sequence 13308, A
455	26	57.8	252	11	US-11-188-298-13728	Sequence 13728, A	528	26	57.8	311	11	US-11-188-298-13308	Sequence 13308, A
456	26	57.8	254	9	US-10-678-556A-177	Sequence 177, App	529	26	57.8	311	11	US-11-188-298-13308	Sequence 13308, A
457	26	57.8	256	11	US-11-188-298-17217	Sequence 17217, A	530	26	57.8	311	11	US-11-188-298-13308	Sequence 13308, A
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535	26	57.8	312	11	US-11-188-298-1843	Sequence 5843, App	608	26	57.8	336	11	US-11-188-298-18889	Sequence 18889, A
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537	26	57.8	313	11	US-11-188-298-17907	Sequence 17907, A	610	26	57.8	336	11	US-11-188-298-21288	Sequence 21288, A
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550	26	57.8	319	9	US-10-454-437-114	Sequence 114, App	623	26	57.8	339	11	US-11-188-298-7093	Sequence 7093, App
551	26	57.8	319	9	US-10-454-437-116	Sequence 116, App	624	26	57.8	339	11	US-11-188-298-21617	Sequence 21617, A
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554	26	57.8	321	11	US-11-188-298-1115	Sequence 1115, App	627	26	57.8	340	11	US-11-188-298-9181	Sequence 9181, App
555	26	57.8	323	11	US-11-096-5688A-8709	Sequence 8709, App	628	26	57.8	340	11	US-11-188-298-114731	Sequence 114731, A
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558	26	57.8	330	11	US-11-188-298-1541	Sequence 1541, App	631	26	57.8	345	9	US-10-467-657-252	Sequence 252, App
559	26	57.8	330	11	US-11-188-298-2592	Sequence 2592, App	632	26	57.8	345	9	US-10-467-657-3086	Sequence 3086, App
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561	26	57.8	330	11	US-11-188-298-16486	Sequence 16486, A	634	26	57.8	347	11	US-11-188-298-1844	Sequence 1844, App
562	26	57.8	332	11	US-11-188-298-3112	Sequence 3112, App	635	26	57.8	347	11	US-11-188-298-19152	Sequence 19152, A
563	26	57.8	332	11	US-11-188-298-4331	Sequence 4331, App	636	26	57.8	350	11	US-11-096-668A-11440	Sequence 11440, App
564	26	57.8	332	11	US-11-188-298-6916	Sequence 8916, App	637	26	57.8	350	11	US-11-096-668A-3576	Sequence 3576, App
565	26	57.8	332	11	US-11-188-298-12219	Sequence 12219, A	638	26	57.8	351	11	US-11-188-298-3576	Sequence 32, App1
566	26	57.8	332	11	US-11-188-298-14513	Sequence 14513, A	639	26	57.8	353	11	US-11-232-367-32	Sequence 32, App1
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568	26	57.8	332	11	US-11-188-298-20533	Sequence 20533, A	641	26	57.8	353	11	US-11-188-298-3401	Sequence 3401, App
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571	26	57.8	333	11	US-11-188-298-29811	Sequence 29811, App	644	26	57.8	356	11	US-10-467-657-8540	Sequence 8540, App
572	26	57.8	333	11	US-11-188-298-5377	Sequence 5377, App	645	26	57.8	357	9	US-11-188-298-8001	Sequence 8001, App
573	26	57.8	333	11	US-11-188-298-7755	Sequence 7755, App	646	26	57.8	359	11	US-11-188-298-9003	Sequence 9003, App
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575	26	57.8	333	11	US-11-188-298-8433	Sequence 8433, App	648	26	57.8	362	11	US-11-087-099-868	Sequence 868, App
576	26	57.8	333	11	US-11-188-298-8597	Sequence 8597, App	649	26	57.8	364	11	US-10-330-773-446	Sequence 446, App
577	26	57.8	333	11	US-11-188-298-11855	Sequence 11855, A	650	26	57.8	374	11	US-11-056-568A-19910	Sequence 19910, A
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582	26	57.8	334	11	US-11-188-298-6304	Sequence 9122, App	655	26	57.8	383	9	US-10-793-628-3110	Sequence 3110, App
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586	26	57.8	335	11	US-11-188-298-733	Sequence 733, App	659	26	57.8	387	11	US-11-188-298-686	Sequence 686, App
587	26	57.8	335	11	US-11-188-298-1070	Sequence 1070, App	660	26	57.8	387	11	US-11-188-298-14095	Sequence 14095, A
588	26	57.8	335	11	US-11-188-298-4651	Sequence 4651, App	661	26	57.8	388	11	US-11-188-298-1638	Sequence 4638, App
589	26	57.8	335	11	US-11-188-298-6345	Sequence 6345, App	662	26	57.8	389	11	US-11-188-298-1989	Sequence 2989, App
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592	26	57.8	335	11	US-11-188-298-13985	Sequence 13985, A	665	26	57.8	391	11	US-11-087-099-4757	Sequence 4757, App
593	26	57.8	335	11	US-11-188-298-14676	Sequence 14676, A	666	26	57.8	392	11	US-11-188-298-21147	Sequence 21147, A
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604	26	57.8	336	11	US-11-188-298-13075	Sequence 13075, A	677	26	57.8	401	11	US-11-055-822-720	Sequence 720, App
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683	26	57.8	406	11	US-11-096-568A-12538	Sequence 12538, A	756	26	57.8	748	11	US-11-087-099-12342	Sequence 12342, A
684	26	57.8	409	11	US-11-096-568A-12537	Sequence 12537, A	757	26	57.8	753	11	US-10-915-002-167	Sequence 167, App
685	26	57.8	414	11	US-11-096-568A-12537	Sequence 12537, A	758	26	57.8	755	9	US-10-131-826A-28	Sequence 28, App1
686	26	57.8	415	11	US-11-096-568A-10224	Sequence 10224, A	759	26	57.8	755	9	US-10-973-1158-28	Sequence 28, App1
687	26	57.8	416	11	US-10-793-626-1406	Sequence 1406, App	760	26	57.8	755	9	US-10-137-873A-28	Sequence 28, App1
688	26	57.8	418	11	US-11-225-709-41	Sequence 41, App1	761	26	57.8	755	9	US-10-152-370-28	Sequence 28, App1
689	26	57.8	419	11	US-11-188-298-2572	Sequence 2572, Ap	762	26	57.8	755	11	US-11-290-153-28	Sequence 28, App1
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691	26	57.8	424	11	US-11-096-568A-25737	Sequence 25737, A	764	26	57.8	810	9	US-10-453-372-1116	Sequence 1116, Ap
692	26	57.8	427	11	US-11-098-686-11271	Sequence 11271, A	765	26	57.8	817	9	US-10-453-372-520	Sequence 520, App
693	26	57.8	429	11	US-11-188-298-10615	Sequence 10615, A	766	26	57.8	833	9	US-10-453-372-519	Sequence 518, App
694	26	57.8	430	11	US-11-096-568A-1439	Sequence 1439, Ap	767	26	57.8	840	9	US-10-915-002-197	Sequence 197, App
695	26	57.8	433	11	US-11-188-298-8416	Sequence 8416, Ap	768	26	57.8	840	9	US-10-915-002-200	Sequence 200, App
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697	26	57.8	439	11	US-11-096-568A-1438	Sequence 1438, Ap	770	26	57.8	858	11	US-11-165-819-5	Sequence 5, App1
698	26	57.8	440	11	US-11-188-298-19715	Sequence 19715, A	771	26	57.8	878	9	US-10-204-639-27	Sequence 27, App1
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702	26	57.8	456	11	US-11-096-568A-12536	Sequence 12536, A	775	26	57.8	898	9	US-10-453-372-522	Sequence 522, App
703	26	57.8	456	11	US-11-096-568A-17833	Sequence 17833, A	776	26	57.8	898	11	US-11-183-136-14	Sequence 14, App1
704	26	57.8	458	9	US-10-934-944-284	Sequence 284, App	777	26	57.8	899	9	US-10-453-372-506	Sequence 506, App
705	26	57.8	458	11	US-11-116-881A-293	Sequence 293, App	778	26	57.8	899	9	US-10-453-372-524	Sequence 524, App
706	26	57.8	464	11	US-11-096-568A-10223	Sequence 10223, A	779	26	57.8	899	9	US-10-453-372-526	Sequence 526, App
707	26	57.8	465	11	US-11-188-298-14054	Sequence 14054, A	780	26	57.8	899	9	US-10-453-372-528	Sequence 528, App
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710	26	57.8	471	11	US-11-045-004-1855	Sequence 1855, Ap	783	26	57.8	899	9	US-10-453-372-534	Sequence 534, App
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715	26	57.8	484	11	US-11-087-099-6811	Sequence 6811, Ap	788	26	57.8	899	9	US-10-453-372-544	Sequence 544, App
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717	26	57.8	485	11	US-11-096-568A-17832	Sequence 17832, A	790	26	57.8	917	11	US-11-169-041-145	Sequence 145, App
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ALIGNMENTS

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RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
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US-10-530-253-13
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Best Local Similarity 100.0%; Pred. No. 0.062;
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RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US2006003919A1
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; GENERAL INFORMATION:
; APPLICANT: HealthBanks Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; CURRENT FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
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US-11-206-138-3
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RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
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Query Match 100.0%; Score 45; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 LQTTIHDI 9
Db 19 LQTTIHDI 27

RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
```


;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 3
;; LENGTH: 248
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 100.0%; Score 45; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 19 LQTTIHDI 27

RESULT 5
US-10-530-253-5
; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 45; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 19 LQTTIHDI 27

RESULT 6
US-10-530-253-7
; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 248

;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 100.0%; Score 45; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 116 LQTTIHDI 124

RESULT 7
US-10-530-253-9
; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 45; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 LQTTIHDI 9
Db 116 LQTTIHDI 124

RESULT 8
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 45; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.11; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 LQTTIHDI 9
Db 116 LQTTIHDI 124

RESULT 9
US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANU, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; CURRENT FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match
Best Local Similarity 100.0%; Score 45; DB 11; Length 256;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 124 LQTTIHDI 132

RESULT 10
US-11-096-568A-1943
; Sequence 1943, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1943
; LENGTH: 250
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(250)
; OTHER INFORMATION: Ceres Seq. ID no. 15180148
US-11-096-568A-1943

Query Match
Best Local Similarity 77.8%; Score 35; DB 11; Length 250;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 7
Db 181 LQTTIHDI 187

RESULT 11
US-11-096-568A-19577

; Sequence 19577, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19577
; LENGTH: 287
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(287)
; OTHER INFORMATION: Ceres Seq. ID no. 12373656
US-11-096-568A-19577

Query Match
Best Local Similarity 77.8%; Score 35; DB 11; Length 287;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
Db 94 VKTTHDIM 102

RESULT 12
US-11-096-568A-1942
; Sequence 1942, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1942
; LENGTH: 298
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (1)-(298)
; OTHER INFORMATION: Ceres Seq. ID no. 15180147
US-11-096-568A-1942

Query Match
Best Local Similarity 77.8%; Score 35; DB 11; Length 298;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 7
Db 229 LQTTIHDI 235

RESULT 13
US-11-096-568A-1941
; Sequence 1941, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 1941

LENGTH: 303
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc.feature
LOCATION: (1)..(303)
OTHER INFORMATION: Ceres Seq. ID no. 15180146
US-11-096-568A-1941

Query Match 77.8%; Score 35; DB 11; Length 303;
Best Local Similarity 85.7%; Pred. No. 14;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTIHDI 7
Db 234 LQTIHDI 240

RESULT 14
US-11-096-568A-6040
; Sequence 6040, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6040
; LENGTH: 329
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(329)
; OTHER INFORMATION: Ceres Seq. ID no. 14313169
US-11-096-568A-6040

Query Match 77.8%; Score 35; DB 11; Length 329;
Best Local Similarity 66.7%; Pred. No. 15;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTIHDI 9
Db 203 LQTIHDI 211

RESULT 15
US-11-096-568A-19576
; Sequence 19576, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19576
; LENGTH: 448
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(448)
; OTHER INFORMATION: Ceres Seq. ID no. 12373655
US-11-096-568A-19576

Query Match 77.8%; Score 35; DB 11; Length 448;
Best Local Similarity 66.7%; Pred. No. 21;

Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTIHDI 9
Db 255 VKTIHDI 263

RESULT 16
US-11-096-568A-19575
; Sequence 19575, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 19575
; LENGTH: 463
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(463)
; OTHER INFORMATION: Ceres Seq. ID no. 12373654
US-11-096-568A-19575

Query Match 77.8%; Score 35; DB 11; Length 463;
Best Local Similarity 66.7%; Pred. No. 22;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 1 LQTIHDI 9
Db 270 VKTIHDI 278

RESULT 17
US-11-096-568A-6039
; Sequence 6039, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6039
; LENGTH: 540
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(540)
; OTHER INFORMATION: Ceres Seq. ID no. 14313168
US-11-096-568A-6039

Query Match 77.8%; Score 35; DB 11; Length 540;
Best Local Similarity 66.7%; Pred. No. 26;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 LQTIHDI 9
Db 414 LQTIHDI 422

RESULT 18
US-11-188-298-5563
; Sequence 5563, Application US/11188298
; Publication No. US20060075522A1

```

; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 5563
; LENGTH: 989
; TYPE: PRT
; ORGANISM: Trichodesmium erythraeum IMS101
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(989)
; OTHER INFORMATION: unsure at all Xaa locations
US-11-188-298-5563
```

```

Query Match          77.8%; Score 35; DB 11; Length 989,
Best Local Similarity 62.5%; Pred. No. 52;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      2 QTTIHDI 9
Db      457 KTTIHDI 464
```

```

RESULT 19
US-10-530-253-17
; Sequence 17, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 17
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 33
US-10-530-253-17
```

```

Query Match          73.3%; Score 33; DB 9; Length 149;
Best Local Similarity 75.0%; Pred. No. 16;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 LOTTIHDI 8
Db      19 LOTTIHDI 26
```

```

RESULT 20
US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
```

```

; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 26
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 68
US-10-530-253-26
```

```

Query Match          73.3%; Score 33; DB 9; Length 158;
Best Local Similarity 62.5%; Pred. No. 17;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LOTTIHDI 8
Db      21 LOTTIHDI 28
```

```

RESULT 21
US-10-506-454-1603
; Sequence 1603, Application US/10506454
; Publication No. US20060068386A1
; GENERAL INFORMATION:
; APPLICANT: Slesarev, Alexi I
; APPLICANT: Mezhevaeva, Katja V
; APPLICANT: Polushin, Nikolai N
; APPLICANT: Shcherbina, Olga V
; APPLICANT: Shakhova, Vera V
; APPLICANT: Mal'kh, Andrei G
; APPLICANT: Kozavkin, Sergei A
; TITLE OF INVENTION: The Complete Genome and Protein Sequences of the Hyperthermophile
; TITLE OF INVENTION: Methanopyrus Kandleri AV19 and Monophyly of Archaeal Methanogens
; FILE REFERENCE: FID001
; CURRENT APPLICATION NUMBER: US/10/506,454
; CURRENT FILING DATE: 2004-08-31
; PRIOR APPLICATION NUMBER: PCT/US03/06664
; PRIOR FILING DATE: 2003-03-04
; PRIOR APPLICATION NUMBER: 60/361,742
; PRIOR FILING DATE: 2002-03-04
; NUMBER OF SEQ ID NOS: 1722
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 1603
; LENGTH: 237
; TYPE: PRT
; ORGANISM: Methanopyrus kandleri
US-10-506-454-1603
```

```

Query Match          73.3%; Score 33; DB 9; Length 237;
Best Local Similarity 62.5%; Pred. No. 27;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LOTTIHDI 8
Db      167 IHTTVHDI 174
```

```

RESULT 22
US-11-188-298-17532
; Sequence 17532, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
```

;; PRIOR FILING DATE: 2004-07-31
;; NUMBER OF SEQ ID NOS: 22569
;; SEQ ID NO 17532
;; LENGTH: 333
;; TYPE: PRT
;; ORGANISM: Magnetococcus sp. MC-1
US-11-188-298-17532

Query Match 71.1%; Score 32; DB 11; Length 333;
Best Local Similarity 100.0%; Pred. No. 62;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTTHDI 8
|||||
Db 175 TTTHDI 180

RESULT 23
US-11-188-298-16137
; Sequence 16137, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16137
; LENGTH: 337
; TYPE: PRT
; ORGANISM: Shewanella oneidensis MR-1
US-11-188-298-16137

Query Match 71.1%; Score 32; DB 11; Length 337;
Best Local Similarity 100.0%; Pred. No. 62;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTTHDI 8
|||||
Db 176 TTTHDI 181

RESULT 24
US-11-188-298-6731
; Sequence 6731, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 6731
; LENGTH: 340
; TYPE: PRT
; ORGANISM: SYNECHOCOCCUS SP. WH 8102
US-11-188-298-6731

Query Match 71.1%; Score 32; DB 11; Length 340;
Best Local Similarity 100.0%; Pred. No. 63;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTTHDI 8
|||||
Db 181 TTTHDI 186

RESULT 25
US-10-873-528-37
; Sequence 37, Application US/10873528
; Publication No. US20050276814A1
; GENERAL INFORMATION:
; APPLICANT: Microbial Technics Limited
; APPLICANT: Gilbert, Christophe FG
; APPLICANT: Hanebro, Philip M
; TITLE OF INVENTION: Proteins
; FILE REFERENCE: PWC/P2129WO
; CURRENT APPLICATION NUMBER: US/10/873,528
; PRIOR FILING DATE: 2004-06-23
; PRIOR APPLICATION NUMBER: US/09/769,787
; PRIOR FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: GB 9816337.1
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: US 60/125164
; PRIOR FILING DATE: 1999-03-19
; NUMBER OF SEQ ID NOS: 388
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 37
; LENGTH: 513
; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-10-873-528-37

Query Match 71.1%; Score 32; DB 9; Length 513;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTTHDI 8
|||||
Db 495 TTTHDI 500

RESULT 26
US-10-873-528-193
; Sequence 193, Application US/10873528
; Publication No. US20050276814A1
; GENERAL INFORMATION:
; APPLICANT: Microbial Technics Limited
; APPLICANT: Gilbert, Christophe FG
; APPLICANT: Hanebro, Philip M
; TITLE OF INVENTION: Proteins
; FILE REFERENCE: PWC/P2129WO
; CURRENT APPLICATION NUMBER: US/10/873,528
; PRIOR FILING DATE: 2004-06-23
; PRIOR APPLICATION NUMBER: US/09/769,787
; PRIOR FILING DATE: 2001-01-26
; PRIOR APPLICATION NUMBER: GB 9816337.1
; PRIOR FILING DATE: 1998-03-27
; PRIOR APPLICATION NUMBER: US 60/125164
; PRIOR FILING DATE: 1999-03-19
; NUMBER OF SEQ ID NOS: 388
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 193
; LENGTH: 513
; TYPE: PRT
; ORGANISM: Streptococcus pneumoniae
US-10-873-528-193

Query Match 71.1%; Score 32; DB 9; Length 513;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTTHDI 8
|||||
Db 495 TTTHDI 500

RESULT 27
US-11-103-957-61
; Sequence 61, Application US/11103957

```

; Publication No. US20050201847A1
; GENERAL INFORMATION:
; APPLICANT: Berthet, Francois-Xavier Jacques
; APPLICANT: Lobet, Yves
; APPLICANT: Poolman, Jan
; APPLICANT: Verlant, Vincent Georges Christian Louis
; TITLE OF INVENTION: Vaccine Composition
; FILE REFERENCE: B45261
; CURRENT APPLICATION NUMBER: US/11/103,957
; PRIOR FILING DATE: 2005-04-12
; PRIOR APPLICATION NUMBER: US/10/467,534
; PRIOR FILING DATE: 2004-02-03
; PRIOR APPLICATION NUMBER: PCT/EP02/01356
; PRIOR FILING DATE: 2002-02-08
; PRIOR APPLICATION NUMBER: GB 0103169.9
; PRIOR FILING DATE: 2001-02-08
; NUMBER OF SEQ ID NOS: 108
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 61
; LENGTH: 553
; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
; US-11-103-957-61

Query Match      71.1%; Score 32; DB 11; Length 553;
Best Local Similarity 75.0%; Pred. No. 1.1e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 LQTTIHDI 8
Db      284 LQTTIHDI 291

RESULT 28
US-11-087-099-1991
; Sequence 1991, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B BP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 1991
; LENGTH: 643
; TYPE: PRT
; ORGANISM: Magnetospirillum magnetotacticum
; US-11-087-099-1991

Query Match      71.1%; Score 32; DB 11; Length 643;
Best Local Similarity 85.7%; Pred. No. 1.3e+02;
Matches 6; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY      1 LQTTIHDI 7
Db      5 LQTTIHDI 11

RESULT 29
US-11-087-099-10391
; Sequence 10391, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B BP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 10391
; LENGTH: 186
; TYPE: PRT

```

```

; ORGANISM: Thermoplasma volcanium
; US-11-087-099-10391

Query Match      68.9%; Score 31; DB 11; Length 186;
Best Local Similarity 44.4%; Pred. No. 51;
Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY      1 LQTTIHDI 9
Db      41 LQTTIHDI 49

RESULT 30
US-11-087-099-7508
; Sequence 7508, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B BP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 7508
; LENGTH: 187
; TYPE: PRT
; ORGANISM: Thermoplasma volcanium
; US-11-087-099-7508

Query Match      68.9%; Score 31; DB 11; Length 187;
Best Local Similarity 44.4%; Pred. No. 51;
Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY      1 LQTTIHDI 9
Db      42 LQTTIHDI 50

RESULT 31
US-11-188-298-21765
; Sequence 21765, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 21765
; LENGTH: 335
; TYPE: PRT
; ORGANISM: Rhodobacter sphaeroides
; US-11-188-298-21765

Query Match      68.9%; Score 31; DB 11; Length 335;
Best Local Similarity 83.3%; Pred. No. 99;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 TTTHDI 8
Db      176 TTTHDI 181

RESULT 32
US-11-188-298-16409
; Sequence 16409, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT

```

```
FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16409
; LENGTH: 340
; TYPE: PRT
; ORGANISM: Prochlorococcus marinus subsp. pastoris str. CCMP1378
US-11-188-298-16409

Query Match      68.9%; Score 31; DB 11; Length 340;
Best Local Similarity 83.3%; Pred. No. 1e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY      3 TTTHDI 8
DB      181 TTTHDV 186

RESULT 33
US-10-703-799B-62
; Sequence 62, Application US/10703799B
; Publication No. US20060078884A1
; GENERAL INFORMATION:
; APPLICANT: Pompejus, Markus
; APPLICANT: Krieger, Burkhard
; APPLICANT: Schröder, Hartwig
; APPLICANT: Zeidler, Oskar
; APPLICANT: Haberdauer, Gregor
; APPLICANT: Lee, Heung-Joon
; APPLICANT: Kim, Hyun-Slick
; TITLE OF INVENTION: CORYNEBACTERIUM GLUTAMICUM GENES ENCODING STRESS,
; TITLE OF INVENTION: RESISTANCE AND TOLERANCE PROTEINS
; FILE REFERENCE: BGI-124CPN
; CURRENT APPLICATION NUMBER: US/10/703,799B
; CURRENT FILING DATE: 2003-11-07
; PRIOR APPLICATION NUMBER: 09/603,208
; PRIOR FILING DATE: 2000-06-23
; PRIOR APPLICATION NUMBER: 60/141031
; PRIOR FILING DATE: 1999-06-25
; PRIOR APPLICATION NUMBER: 60/142692
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: 60/151214
; PRIOR FILING DATE: 1999-08-27
; PRIOR APPLICATION NUMBER: DE 19930429.7
; PRIOR FILING DATE: 1999-07-01
; PRIOR APPLICATION NUMBER: DE 19931413.6
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19931457.8
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19931541.8
; PRIOR FILING DATE: 1999-07-08
; PRIOR APPLICATION NUMBER: DE 19932209.0
; PRIOR FILING DATE: 1999-07-09
; PRIOR APPLICATION NUMBER: DE 19932230.9
; PRIOR FILING DATE: 1999-07-09
; Remaining Prior Application data removed - See file wrapper or PALM.
; NUMBER OF SEQ ID NOS: 306
; SEQ ID NO 62
; LENGTH: 383
; TYPE: PRT
; ORGANISM: Corynebacterium glutamicum
US-10-703-799B-62

Query Match      68.9%; Score 31; DB 9; Length 383;
Best Local Similarity 55.6%; Pred. No. 1.1e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LQTTIHDI 9
DB      238 MKKTTIHDI 246

RESULT 34
US-10-503-575-174
; Sequence 174, Application US/10503575
; Publication No. US20050244823A1
; GENERAL INFORMATION:
; APPLICANT: Drijfhout, Jan Wouter
; APPLICANT: van Veele, Petrus Antonius
; APPLICANT: Konig, Frits
; TITLE OF INVENTION: NOVEL EPITOPES FOR CELIAC DISEASE AND AUTOIMMUNE DISEASES, METHOD
; TITLE OF INVENTION: DETECTING THOSE AND NOVEL NON-ANTIGENIC FOOD COMPOUNDS
; FILE REFERENCE: 2799/72843-PCT-US
; CURRENT APPLICATION NUMBER: US/10/503,575
; CURRENT FILING DATE: 2004-08-04
; PRIOR APPLICATION NUMBER: PCT/NL03/00077
; PRIOR FILING DATE: 2003-02-04
; PRIOR APPLICATION NUMBER: EP 02075456.0
; PRIOR FILING DATE: 2002-02-04
; NUMBER OF SEQ ID NOS: 340
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 174
; LENGTH: 19
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-503-575-174

Query Match      66.7%; Score 30; DB 9; Length 19;
Best Local Similarity 71.4%; Pred. No. 6.3;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY      1 LQTTIHDI 7
DB      9 LQATIHDI 15

RESULT 35
US-11-079-463-6103
; Sequence 6103, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRA
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PTH00-03DIY2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6103
; LENGTH: 63
; TYPE: PRT
; ORGANISM: B. fragilis
US-11-079-463-6103

Query Match      66.7%; Score 30; DB 11; Length 63;
Best Local Similarity 62.5%; Pred. No. 24;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 LQTTIHDI 8
DB      16 IQASTIHDI 23

RESULT 36
US-10-986-405-199
; Sequence 199, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
```

```
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; CURRENT FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 199
; LENGTH: 133
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-199
```

```
Query Match          66.7%; Score 30; DB 9; Length 133;
Best Local Similarity 71.4%; Pred. No. 56;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LQTTIHD 7
        |||:|
Db      107 LQATLHD 113
```

```
RESULT 37
US-10-986-405-205
; Sequence 205, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; CURRENT FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 205
; LENGTH: 135
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-205
```

```
Query Match          66.7%; Score 30; DB 9; Length 135;
Best Local Similarity 71.4%; Pred. No. 57;
```

```
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
QY      1 LQTTIHD 7
        |||:|
Db      107 LQATLHD 113
```

```
RESULT 38
US-10-986-405-229
; Sequence 229, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; CURRENT FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
; PRIOR FILING DATE: 2002-10-04
; PRIOR APPLICATION NUMBER: US 60/402,799
; PRIOR FILING DATE: 2002-08-13
; NUMBER OF SEQ ID NOS: 378
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 229
; LENGTH: 140
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-986-405-229
```

```
Query Match          66.7%; Score 30; DB 9; Length 140;
Best Local Similarity 71.4%; Pred. No. 59;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 LQTTIHD 7
        |||:|
Db      122 LQATLHD 128
```

```
RESULT 39
US-10-986-405-200
; Sequence 200, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
; CURRENT FILING DATE: 2004-11-12
; PRIOR APPLICATION NUMBER: PCT/US2003/015439
; PRIOR FILING DATE: 2003-05-16
; PRIOR APPLICATION NUMBER: US 60/388,543
; PRIOR FILING DATE: 2002-06-12
; PRIOR APPLICATION NUMBER: US 60/401,757
; PRIOR FILING DATE: 2002-08-08
; PRIOR APPLICATION NUMBER: US 60/381,592
; PRIOR FILING DATE: 2002-05-17
; PRIOR APPLICATION NUMBER: US 60/402,585
; PRIOR FILING DATE: 2002-08-12
; PRIOR APPLICATION NUMBER: US 60/404,959
; PRIOR FILING DATE: 2002-08-22
; PRIOR APPLICATION NUMBER: US 60/415,902
```



```
;; PRIOR FILING DATE: 2002-10-04
;; PRIOR APPLICATION NUMBER: US 60/402,799
;; PRIOR FILING DATE: 2002-08-13
;; NUMBER OF SEQ ID NOS: 378
;; SOFTWARE: PatentIn Ver. 2.0
;; SEQ ID NO 200
;; LENGTH: 149
;; TYPE: PRT
;; ORGANISM: Homo sapiens
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (50)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (122)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (138)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (143)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-10-986-405-200
```

```
Query Match          66.7%; Score 30; DB 9; Length 149;
Best Local Similarity 71.4%; Pred. No. 63;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 1QTTIHD 7
        |||:|
DB      107 1QATLHD 113
```

```
RESULT 40
US-11-188-298-10295
; Sequence 10295, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 10295
; LENGTH: 179
; TYPE: PRT
; ORGANISM: Methanothermobacter thermoautotrophicus str. Delta H
US-11-188-298-10295
```

```
Query Match          66.7%; Score 30; DB 11; Length 179;
Best Local Similarity 66.7%; Pred. No. 78;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 1QTTIHD 11 9
        |||:|
DB      74 1LISTIHD 82
```

```
RESULT 41
US-10-986-405-216
; Sequence 216, Application US/10986405
; Publication No. US20060073561A1
; GENERAL INFORMATION:
; APPLICANT: Rosen, et al.
; TITLE OF INVENTION: 157 Human Secreted Proteins
; FILE REFERENCE: P2750P1
; CURRENT APPLICATION NUMBER: US/10/986,405
```

```
;; CURRENT FILING DATE: 2004-11-12
;; PRIOR APPLICATION NUMBER: PCT/US2003/015439
;; PRIOR FILING DATE: 2003-05-16
;; PRIOR APPLICATION NUMBER: US 60/388,543
;; PRIOR FILING DATE: 2002-06-12
;; PRIOR APPLICATION NUMBER: US 60/401,757
;; PRIOR FILING DATE: 2002-08-08
;; PRIOR APPLICATION NUMBER: US 60/381,592
;; PRIOR FILING DATE: 2002-05-17
;; PRIOR APPLICATION NUMBER: US 60/402,585
;; PRIOR FILING DATE: 2002-08-12
;; PRIOR APPLICATION NUMBER: US 60/404,959
;; PRIOR FILING DATE: 2002-08-22
;; PRIOR APPLICATION NUMBER: US 60/415,902
;; PRIOR FILING DATE: 2002-10-04
;; PRIOR APPLICATION NUMBER: US 60/402,799
;; PRIOR FILING DATE: 2002-08-13
;; NUMBER OF SEQ ID NOS: 378
;; SOFTWARE: PatentIn Ver. 2.0
;; SEQ ID NO 216
;; LENGTH: 184
;; TYPE: PRT
;; ORGANISM: Homo sapiens
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (9)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (50)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (98)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (137)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
;; FEATURE:
;; NAME/KEY: SITE
;; LOCATION: (163)
;; OTHER INFORMATION: Xaa equals any of the naturally occurring L-amino acids
US-10-986-405-216
```

```
Query Match          66.7%; Score 30; DB 9; Length 184;
Best Local Similarity 71.4%; Pred. No. 80;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 1QTTIHD 7
        |||:|
DB      107 1QATLHD 113
```

```
RESULT 42
US-11-188-298-17244
; Sequence 17244, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 17244
; LENGTH: 291
; TYPE: PRT
; ORGANISM: Streptomyces galliaueus
US-11-188-298-17244
```

Query Match 66.7%; Score 30; DB 11; Length 291;
Best Local Similarity 50.0%; Pred. No. 1.3e+02;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 LOTTIHI 8
||:||||:
Db 120 LQSNVHDV 127

RESULT 43
US-11-188-298-3948

; Sequence 3948, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 3948
; LENGTH: 333
; TYPE: PRT
; ORGANISM: Vibrio vulnificus CMCP6

US-11-188-298-3948

Query Match 66.7%; Score 30; DB 11; Length 333;
Best Local Similarity 83.3%; Pred. No. 1.6e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTIHDI 8
|||||:
Db 176 TTIHDL 181

RESULT 44
US-11-188-298-22053

; Sequence 22053, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 22053
; LENGTH: 333
; TYPE: PRT
; ORGANISM: Vibrio parahaemolyticus RIMD 2210633

US-11-188-298-22053

Query Match 66.7%; Score 30; DB 11; Length 333;
Best Local Similarity 83.3%; Pred. No. 1.6e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTIHDI 8
|||||:
Db 176 TTIHDL 181

RESULT 45
US-11-188-298-2051

; Sequence 2051, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B

; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 2051
; LENGTH: 334
; TYPE: PRT
; ORGANISM: Azotobacter vinelandii

US-11-188-298-2051

Query Match 66.7%; Score 30; DB 11; Length 334;
Best Local Similarity 83.3%; Pred. No. 1.6e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 TTIHDI 8
|||||:
Db 176 TTIHDL 181

RESULT 46
US-11-096-568A-21865

; Sequence 21865, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 21865
; LENGTH: 338
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(338)
; OTHER INFORMATION: Cereals Seq. ID no. 12406582

US-11-096-568A-21865

Query Match 66.7%; Score 30; DB 11; Length 338;
Best Local Similarity 57.1%; Pred. No. 1.6e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 QTTIHDI 8
||:||||:
Db 295 ETTIHVDV 301

RESULT 47
US-11-188-298-12429

; Sequence 12429, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; PRIOR FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 12429
; LENGTH: 352
; TYPE: PRT
; ORGANISM: Rhodococcus sp. Phi2

US-11-188-298-12429

Query Match 66.7%; Score 30; DB 11; Length 352;
Best Local Similarity 83.3%; Pred. No. 1.7e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 TTTHDI 8
| | | | |
Db 11 LTRVHSIL 16

RESULT 48

US-10-878-556A-84
; Sequence 84, Application US/10878556A
; Publication No. US2005026399A1
; GENERAL INFORMATION:
; APPLICANT: Hoffmann La-Roche Inc.
; TITLE OF INVENTION: HCV regulated protein expression
; FILE REFERENCE: 21762
; CURRENT APPLICATION NUMBER: US/10/878,556A
; CURRENT FILING DATE: 2004-06-28
; NUMBER OF SEQ ID NOS: 199
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 84
; LENGTH: 354
; TYPE: PRT
; ORGANISM: Homo sapiens
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: sw_hum/arg2_human
; DATABASE ENTRY DATE: 1997-11-01
US-10-878-556A-84

Query Match 66.7%; Score 30; DB 9; Length 354;
Best Local Similarity 55.6%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
| | | | |
Db 11 LQTRVHSIL 19

RESULT 49

US-11-100-640-4
; Sequence 4, Application US/11100640
; Publication No. US2006003524A1
; GENERAL INFORMATION:
; APPLICANT: Riggs, Gregory J
; TITLE OF INVENTION: A Method for Distinguishing Follicular Thyroid Adenoma (FTA) from
; TITLE OF INVENTION: Follicular Thyroid Carcinoma (FTC)
; FILE REFERENCE: 000250.00021
; CURRENT APPLICATION NUMBER: US/11/100,640
; CURRENT FILING DATE: 2005-04-07
; NUMBER OF SEQ ID NOS: 98
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 4
; LENGTH: 354
; TYPE: PRT
; ORGANISM: human
US-11-100-640-4

Query Match 66.7%; Score 30; DB 11; Length 354;
Best Local Similarity 55.6%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 LQTTIHDI 9
| | | | |
Db 11 LQTRVHSIL 19

RESULT 50

US-11-079-463-6883
; Sequence 6883, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRA
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: PATH00-03D1V2

;; CURRENT APPLICATION NUMBER: US/11/079,463
;; CURRENT FILING DATE: 2005-03-14
;; PRIOR APPLICATION NUMBER: US 60/128,705
;; PRIOR FILING DATE: 1999-04-09
;; PRIOR APPLICATION NUMBER: US 09/540,209
;; PRIOR FILING DATE: 2000-04-04
;; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 6883
; LENGTH: 371
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-6883

Query Match 66.7%; Score 30; DB 11; Length 371;
Best Local Similarity 83.3%; Pred. No. 1.8e+02;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 3 TTTHDI 8
| | | | |
Db 238 TTTHDI 243

Search completed: May 5, 2006, 08:07:53
Job time : 10.5 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 03:13:35 ; Search time 21 Seconds
(without alignments)
35.432 Million cell updates/sec

Title: US-08-170-344-7
Perfect score: 49
Sequence: 1 T1HDILLEC 9

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Gapop 10.0 , Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

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3: /cgn2_6/ptodata/1/1aa/H-COMB.pep: *
4: /cgn2_6/ptodata/1/1aa/BCTUS-COMB.pep: *
5: /cgn2_6/ptodata/1/1aa/RE-COMB.pep: *
6: /cgn2_6/ptodata/1/1aa/backfile1.pep: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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6	49	100.0	30 2 US-09-980-523A-4	Sequence 4, App1
7	49	100.0	59 2 US-09-390-027-6	Sequence 6, App1
8	49	100.0	151 2 US-09-701-080C-18	Sequence 18, App1
9	49	100.0	158 2 US-09-980-523A-2	Sequence 2, App1
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11	49	100.0	162 1 US-08-316-239B-4	Sequence 4, App1
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13	49	100.0	172 2 US-09-359-382-14	Sequence 14, App1
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15	49	100.0	243 2 US-09-462-993-1	Sequence 10, App1
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17	49	100.0	266 2 US-09-359-382-10	Sequence 10, App1
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22	49	100.0	390 2 US-09-485-885-14	Sequence 14, App1
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25	36	73.5	1164 2 US-09-457-708-2	Sequence 2, App1
26	36	73.5	1164 2 US-09-950-046A-2	Sequence 2, App1
27	36	73.5	1164 2 US-09-976-594-989	Sequence 989, App

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29	34	69.4	111 2 US-09-270-767-33608	Sequence 33608, A
30	34	69.4	111 2 US-09-270-767-48825	Sequence 48825, A
31	34	69.4	266 2 US-09-489-039A-13381	Sequence 13381, A
32	33	67.3	374 2 US-09-638-937-2	Sequence 2, App1
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37	32	65.3	10 2 US-08-159-339A-573	Sequence 60201, A
38	32	65.3	217 2 US-09-270-767-60201	Sequence 29181, A
39	32	65.3	227 2 US-09-252-991A-29181	Sequence 5486, App
40	32	65.3	305 2 US-09-107-532A-5486	Sequence 112, App
41	32	65.3	343 2 US-09-266-965-112	Sequence 22434, A
42	32	65.3	343 2 US-09-248-796A-22434	Sequence 44745, A
43	32	65.3	361 2 US-09-270-767-44745	Sequence 15186, A
44	32	65.3	512 2 US-09-248-796A-15186	Sequence 2, App1
45	32	65.3	520 2 US-09-527-073-2	Sequence 11, App1
46	32	65.3	780 1 US-08-448-196A-9	Sequence 11498, A
47	32	65.3	780 2 US-09-785-381-11	Sequence 15476, A
48	32	65.3	803 2 US-09-949-016-11498	Sequence 102, App
49	32	65.3	1060 2 US-08-787-547-102	Sequence 248, App
50	31	63.3	9 1 US-08-159-339A-248	Sequence 2128, A
51	31	63.3	9 2 US-09-248-796A-21728	Patent No. 5284931
52	31	63.3	66 2 US-08-015-985-13	Sequence 2, App1
53	31	63.3	90 6 5284931-14	Sequence 2, App1
54	31	63.3	139 2 US-09-027-381-2	Sequence 7521, App
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58	31	63.3	235 1 US-08-015-985-13	Sequence 7, App1
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61	31	63.3	254 2 US-09-591-435-7	Sequence 7, App1
62	31	63.3	254 2 US-09-919-039-93	Sequence 42315, A
63	31	63.3	257 2 US-10-098-600B-7	Sequence 14, App1
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65	31	63.3	274 2 US-09-790-045-14	Sequence 14, App1
66	31	63.3	274 2 US-10-222-577-14	Sequence 14, App1
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70	31	63.3	277 2 US-09-248-796A-21807	Sequence 21807, A
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72	31	63.3	297 2 US-09-489-039A-7396	Sequence 7396, App
73	31	63.3	320 2 US-09-949-016-11458	Sequence 4168, App
74	31	63.3	333 2 US-09-107-532A-4769	Sequence 21612, A
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79	31	63.3	372 2 US-09-949-016-11420	Sequence 11420, A
80	31	63.3	412 2 US-09-107-532A-5313	Sequence 5313, App
81	31	63.3	454 2 US-09-489-039A-8358	Sequence 8358, App
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107	30	61.2	142	2	US-08-906-613-155	Sequence 155, App
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118	30	61.2	258	1	US-09-964-956-75	Sequence 75, Appl
119	30	61.2	259	1	US-07-857-224B-49	Sequence 49, Appl
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129	30	61.2	271	2	US-09-540-236-2951	Sequence 2951, App
130	30	61.2	273	1	US-08-701-191A-32	Sequence 32, Appl
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137	30	61.2	336	2	US-09-710-279-3210	Sequence 3210, App
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174	30	61.2	558	2	US-09-252-991A-21305	Sequence 21305, A
175	30	61.2	636	2	US-10-104-047-2449	Sequence 2449, App
176	30	61.2	656	2	US-09-327-984A-36	Sequence 36, Appl
177	30	61.2	777	2	US-09-949-016-1158	Sequence 1158, App
178	30	61.2	931	2	US-09-949-016-9988	Sequence 9988, App
179	30	61.2	943	1	US-08-808-982-7	Sequence 7, Appl
180	30	61.2	943	2	US-09-306-902A-7	Sequence 7, Appl
181	30	61.2	945	2	US-10-037-417-38	Sequence 38, Appl
182	30	61.2	945	2	US-10-037-417-121	Sequence 121, App
183	30	61.2	1075	2	US-09-252-991A-17124	Sequence 17124, A
184	30	61.2	1075	2	US-09-949-016-8308	Sequence 8308, App
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186	30	61.2	1111	2	US-08-800-593-15	Sequence 15, Appl
187	30	61.2	1111	2	US-09-756-071B-15	Sequence 15, Appl
188	30	61.2	1143	2	US-10-104-047-2802	Sequence 2802, App
189	30	61.2	1172	2	US-09-560-385A-28	Sequence 28, Appl
190	30	61.2	1172	2	US-09-560-385A-32	Sequence 32, Appl
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192	30	61.2	1193	1	US-08-317-450B-13	Sequence 13, Appl
193	30	61.2	1193	2	US-08-800-593-13	Sequence 13, Appl
194	30	61.2	1193	2	US-09-560-385A-26	Sequence 26, Appl
195	30	61.2	1193	2	US-09-560-385A-30	Sequence 30, Appl
196	30	61.2	1193	2	US-10-053-662A-31	Sequence 31, Appl
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208	30	61.2	1452	1	US-08-769-399-4	Sequence 4, Appl
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223	30	61.2	3056	2	US-08-952-127-3	Sequence 3, Appl
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278	29	59.2	440	2	US-08-061-062A-8	Sequence 8, Appli	351	28	57.1	106	2	US-10-058-993-90	Sequence 90, Appl
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292	29	59.2	615	1	US-08-023-610-17	Sequence 17, Appl	365	28	57.1	178	2	US-09-267-963D-33	Sequence 23886, A
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299	29	59.2	684	2	US-09-564-805-233	Sequence 233, App	372	28	57.1	219	2	US-10-366-688-2	Sequence 2, Appli
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306	29	59.2	784	1	US-09-184-937-2	Sequence 2, Appli	379	28	57.1	243	2	US-09-902-540-16025	Sequence 16025, A
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408	28	57.1	313	2	US-08-472-679H-192	Sequence 192, App
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433	28	57.1	389	2	US-08-472-679H-114	Sequence 114, App
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468	28	57.1	498	2	US-09-949-016-7712	Sequence 9712, App
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476	28	57.1	512	2	US-09-603-311-24	Sequence 24, App1
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573	28	57.1	970	1	US-08-449-645A-11	Sequence 11, Appl1	647	28	57.1	2628	2	PCT-US94-00198-3	Sequence 3, Appl1
574	28	57.1	970	4	PCT-US95-04681-11	Sequence 11, Appl1	648	28	57.1	3025	6	5223423-3	Patent No. 5223423
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576	28	57.1	984	2	US-08-673-789-6	Sequence 6502, Ap	650	28	57.1	3177	1	US-08-477-451-4	Sequence 4, Appl1
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712	27	55.1	107	2	US-09-621-976-3893	Sequence 3893, Ap	785	27	55.1	235	2	US-09-248-796A-23106	Sequence 45, Appl
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ALIGNMENTS

RESULT 1
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; Patent No. 6419931
; GENERAL INFORMATION:
; APPLICANT: VITHELLO, Maria A.
; APPLICANT: CHESTNUT, Robert W.
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; APPLICANT: CELIS, Etebean
; APPLICANT: GRAY, Howard
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
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; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
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; INFORMATION FOR SEQ ID NO: 72:
; SEQUENCE CHARACTERISTICS:
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Query Match 100.0%; Score 49; DB 2; Length 10;
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Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

RESULT 2
US-09-601-729-273
; Sequence 273, Application US/09601729
; Patent No. 6683052
; GENERAL INFORMATION:
; APPLICANT: THIAM, KADER
; APPLICANT: AURIAULT, CLAUDE
; APPLICANT: GRAS-MASSE, HELENE
; APPLICANT: LOING, ESTELLE
; APPLICANT: VERMAERDE, CLAUDE
; APPLICANT: GUILLET, JEAN GERARD
; TITLE OF INVENTION: LIPOPEPTIDES CONTAINING AN INTERFERON FRAGMENT AND USES
; TITLE OF INVENTION: THEREOF IN PHARMACEUTICAL COMPOSITIONS
; FILE REFERENCE: USB-97-AU-IN
; CURRENT APPLICATION NUMBER: US/09/601,729
; PRIOR FILING DATE: 2000-11-20
; PRIOR APPLICATION NUMBER: PCT/FR99/00259
; PRIOR FILING DATE: 1999-02-05
; PRIOR APPLICATION NUMBER: 98 01439
; PRIOR FILING DATE: 1998-02-06
; NUMBER OF SEQ ID NOS: 281
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 273
; LENGTH: 10
; TYPE: PPT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-601-729-273
Query Match 100.0%; Score 49; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.0046;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 TTHDIIIEC 9
DB 1 TTHDIIIEC 9
RESULT 3
PCT-US95-02121-72
; Sequence 72, Application PC/TUS9502121
; GENERAL INFORMATION:
; APPLICANT:
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
; TITLE OF INVENTION: CTL IMMUNITY
; NUMBER OF SEQUENCES: 153
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/02121
FILING DATE: 16-FEB-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/197,484
FILING DATE: 16-FEB-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935,811
FILING DATE: 26-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,682
FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W.
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4PC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 467-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
PCT-US95-02121-72

Query Match 100.0%; Score 49; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.0046;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD11EC 9
Db 1 T1HD11EC 9

RESULT 4
US-07-909-122-3
Sequence 3, Application US/07909122
Patent No. 5415995
GENERAL INFORMATION:
APPLICANT: SCHOOLNIK, GARY K.
APPLICANT: PALEFSKY, JOEL M.
TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
TITLE OF INVENTION: VIRUS
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/909,122
FILING DATE: 19920706
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:

NAME: BENZ, WILLIAM H.
REGISTRATION NUMBER: 25,952
REFERENCE/DOCKET NUMBER: 28600-20105.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 706141
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 14 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
TOPOLOGY: linear
US-07-909-122-3

Query Match 100.0%; Score 49; DB 1; Length 14;
Best Local Similarity 100.0%; Pred. No. 0.0065;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD11EC 9
Db 6 T1HD11EC 14

RESULT 5
US-08-363-586-4
Sequence 4, Application US/08363586
Patent No. 5629161
GENERAL INFORMATION:
APPLICANT: Mueller, Martin
APPLICANT: Glasmann, Iutz
TITLE OF INVENTION: Use of HPV-16 E6 and E7-Gene Derived
TITLE OF INVENTION: Peptides for the Diagnostic Purpose
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finegan, Henderson, Farabow, Garrett &
ADDRESSER: Dunner
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/363,586
FILING DATE: 23-DEC-1994
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/909,296
FILING DATE: 09-JUL-1992
APPLICATION NUMBER: EP 9111720.8
FILING DATE: 13-JUL-1991
ATTORNEY/AGENT INFORMATION:
NAME: Madler, Linda A.
REGISTRATION NUMBER: 33,218
REFERENCE/DOCKET NUMBER: 02481-1195-00000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-408-4000
TELEFAX: 202-408-4400
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 30 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-363-586-4

Query Match 100.0%; Score 49; DB 1; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.014;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 22 T1HDIILEC 30

RESULT 6
US-09-980-523A-4
; Sequence 4, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-4

Query Match 100.0%; Score 49; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.014;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 15 T1HDIILEC 23

RESULT 7
US-09-390-027-6
; Sequence 6, Application US/09390027
; Patent No. 6235523
; GENERAL INFORMATION:
; APPLICANT: GAJEWICZYK, Diane M.
; APPLICANT: PERSSON, ROY
; APPLICANT: YAO, Fei-Long
; APPLICANT: KLEIN, Michel H.
; APPLICANT: TARTAGLIA, James
; APPLICANT: MOJNGEON, Philippe
; APPLICANT: KOVINSKI, Benjamin
; TITLE OF INVENTION: TREATMENT OF CERVICAL CANCER
; FILE REFERENCE: 1038-982 MIS:jb
; CURRENT APPLICATION NUMBER: US/09/390,027
; PRIOR FILING DATE: 1998-09-03
; EARLIER APPLICATION NUMBER: 60/099,291
; EARLIER FILING DATE: 1998-09-04
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 6
; LENGTH: 59
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-390-027-6

Query Match 100.0%; Score 49; DB 2; Length 59;
Best Local Similarity 100.0%; Pred. No. 0.029;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 50 T1HDIILEC 58

RESULT 8
US-09-701-080C-18
; Sequence 18, Application US/09701080C
; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CRIB BINDING PROTEIN AND RELATED PROTEIN P300 FC
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; PRIOR FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match 100.0%; Score 49; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.077;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 22 T1HDIILEC 30

RESULT 9
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: MOBI AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patentin Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match 100.0%; Score 49; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.081;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
|||
Db 29 TTHDIIIEC 37

RESULT 10
US-08-316-239B-3
; Sequence 3, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHETICAL: NO
US-08-316-239B-3

Query Match 100.0%; Score 49; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.083; 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 TTHDIIIEC 9
|||
Db 29 TTHDIIIEC 37

RESULT 11
US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates

STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtiani, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-4

Query Match 100.0%; Score 49; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.083; 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 TTHDIIIEC 9
|||
Db 29 TTHDIIIEC 37

RESULT 12
US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WREB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14

Query Match 100.0%; Score 49; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.089; 0; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY 1 TTHDIIIEC 9
|||

Db 98 T1HD11EC 106

RESULT 13

US-09-359-382-14

Sequence 14, Application US/09359382

Patent No. 6306397

GENERAL INFORMATION:

APPLICANT: EDWARDS, Stirling John

APPLICANT: COX, John Cooper

APPLICANT: WEBB, Elizabeth Ann

APPLICANT: FRAZER, Ian

TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS

FILE REFERENCE: 017227/0148

CURRENT APPLICATION NUMBER: US/09/359,382

CURRENT FILING DATE: 1999-07-23

EARLIER APPLICATION NUMBER: US 08/860,165

EARLIER FILING DATE: 1997-09-22

EARLIER APPLICATION NUMBER: PCT/AU95/00868

EARLIER FILING DATE: 1995-12-20

EARLIER APPLICATION NUMBER: AU PNO157/94

EARLIER FILING DATE: 1994-12-20

NUMBER OF SEQ ID NOS: 27

SOFTWARE: PatentIn Ver. 2.0

SEQ ID NO 14

LENGTH: 172

TYPE: PRT

ORGANISM: Human papillomavirus type 16

US-09-359-382-14

Query Match 100.0%; Score 49; DB 2; Length 172;

Best Local Similarity 100.0%; Pred. No. 0.089;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HD11EC 9

Db 98 T1HD11EC 106

RESULT 14

US-08-117-083-10

Sequence 10, Application US/08117083

Patent No. 5719054

GENERAL INFORMATION:

APPLICANT: Boutsenell, Michael E.

APPLICANT: Ingile, Stephen C.

APPLICANT: Munro, Alan J.

TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human

TITLE OF INVENTION: Papilloma Virus Proteins

NUMBER OF SEQUENCES: 70

CORRESPONDENCE ADDRESS:

ADDRESSEE: Walter H. Dreyer

STREET: 4 Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: CA

COUNTRY: USA

ZIP: 94111

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/117,083

FILING DATE: 10-SEP-1993

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: Dreyer, Walter H.

REGISTRATION NUMBER: 24,190

REFERENCE/DOCKET NUMBER: A-58783

TELECOMMUNICATION INFORMATION:

TELEPHONE: 415-781-1989

TELEFAX: 415-398-3249

TELEX: 910 277299

INFORMATION FOR SEQ ID NO: 10:

SEQUENCE CHARACTERISTICS:

LENGTH: 182 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: protein

FEATURE:

NAME/KEY: Protein

LOCATION: 1..182

OTHER INFORMATION: /note="Xaa refers to stop codon in

OTHER INFORMATION: the open reading frame."

US-08-117-083-10

Query Match 100.0%; Score 49; DB 1; Length 182;

Best Local Similarity 100.0%; Pred. No. 0.094;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HD11EC 9

Db 30 T1HD11EC 38

RESULT 15

US-09-462-993-1

Sequence 1, Application US/09462993

Patent No. 6884786

GENERAL INFORMATION:

APPLICANT: KIENY, Marie-Paule

APPLICANT: BALLOU, Jean-Marc

APPLICANT: BIRGOURNE, Nadine

TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC

TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION

FILE REFERENCE: 017753-122

CURRENT APPLICATION NUMBER: US/09/462,993

CURRENT FILING DATE: 2000-04-17

PRIOR APPLICATION NUMBER: PCT/FR98/01576

PRIOR FILING DATE: 1998-07-17

PRIOR APPLICATION NUMBER: FR 97/09152

PRIOR FILING DATE: 1997-07-18

NUMBER OF SEQ ID NOS: 23

SOFTWARE: PatentIn Ver. 2.2

SEQ ID NO 1

LENGTH: 243

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: Description of Artificial Sequence: Derived from

OTHER INFORMATION: human papillomavirus, strain HPV-16, B6 protein

OTHER INFORMATION: fused F protein signals, clone B6*TMF.

US-09-462-993-1

Query Match 100.0%; Score 49; DB 2; Length 243;

Best Local Similarity 100.0%; Pred. No. 0.13;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HD11EC 9

Db 57 T1HD11EC 65

RESULT 16

US-08-860-165-10

Sequence 10, Application US/08860165A

Patent No. 6004557

GENERAL INFORMATION:

APPLICANT: EDWARDS, Stirling John

APPLICANT: COX, John Cooper

APPLICANT: WEBB, Elizabeth Ann

APPLICANT: FRAZER, Ian

TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS

FILE REFERENCE: 17227/130

CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P0157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10

Query Match 100.0%; Score 49; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDI1LEC 9
Db 29 T1HDI1LEC 37

RESULT 17
US-09-359-382-10
Sequence 10, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: PEAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P0157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 49; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDI1LEC 9
Db 29 T1HDI1LEC 37

RESULT 18
US-09-367-309A-1
Sequence 1, Application US/09367309A
Patent No. 6428807
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080

PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 49; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDI1LEC 9
Db 29 T1HDI1LEC 37

RESULT 19
US-09-485-885-4
Sequence 4, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 49; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDI1LEC 9
Db 135 T1HDI1LEC 143

RESULT 20
US-09-485-885-10
Sequence 10, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22

NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 49; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 THDIIIEC 9
Db 154 THDIIIEC 162

RESULT 21
US-09-485-885-6
; Sequence 6, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelsh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 49; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 THDIIIEC 9
Db 135 THDIIIEC 143

RESULT 22
US-09-485-885-14
; Sequence 14, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelsh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14

LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 49; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 THDIIIEC 9
Db 154 THDIIIEC 162

RESULT 23
US-08-159-339A-73
; Sequence 73, Application US/08159339A
; Patent No. 6037135
; GENERAL INFORMATION:
; APPLICANT: Kubo, Ralph T.
; APPLICANT: Grey, Howard W.
; APPLICANT: Sette, Alessandro
; APPLICANT: Celis, Esteban
; TITLE OF INVENTION: HLA Binding peptides and Their
; NUMBER OF SEQUENCES: 1254
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew LLP
; STREET: Two Embarcadero Center, Eighth Floor
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111-3834
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/159,339A
; FILING DATE: 29-NOV-1993
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/926,666
; FILING DATE: 07-AUG-1992
; APPLICATION NUMBER: US 08/027,746
; FILING DATE: 05-MAR-1993
; APPLICATION NUMBER: US 08/103,396
; FILING DATE: 06-AUG-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Weber, Ellen Lauer
; REGISTRATION NUMBER: 32,762
; REFERENCE/DOCKET NUMBER: 018623-005030US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 576-0200
; TELEFAX: (415) 576-0300
; TELEX:
; INFORMATION FOR SEQ ID NO: 73:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 10 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-159-339A-73

Query Match 89.8%; Score 44; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 2 THDIIIEC 9
Db 1 THDIIIEC 8

RESULT 24
US-09-248-796A-15013
Sequence 15013, Application US/09248796A
Patent No. 6747137
GENERAL INFORMATION:
APPLICANT: Keith Weinstein et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT FILING DATE: US/09/248,796A
PRIOR FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 15013
LENGTH: 620
TYPE: PRT
ORGANISM: Candida albicans
US-09-248-796A-15013

Query Match
Best Local Similarity 85.7%; Score 42; DB 2; Length 620;
Best Local Similarity 66.7%; Pred. No. 7.1;
Matches 6; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TTHDILLC 9
Db 492 SLHDIVLEC 500

RESULT 25
US-09-457-708-2
Sequence 2, Application US/09457708
Patent No. 6326483
GENERAL INFORMATION:
APPLICANT: Kwiatkowski, David J.
APPLICANT: Sampson, Julian R.
APPLICANT: Povey, Sue
APPLICANT: van Slegtenhorst, Marjon
APPLICANT: Halley, Dicky
TITLE OF INVENTION: Compositions and Methods Based U
TITLE OF INVENTION: Sclerosis-1 (TSC-1) Gene and Gene Product
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSER: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Releasee #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/457,708
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/42002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6604
TELEFAX: (202) 639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-457-708-2

Query Match
Best Local Similarity 73.5%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIILLC 9
Db 253 HDVIVLC 259

RESULT 26
US-09-950-046A-2
Sequence 2, Application US/09950046A
Patent No. 6548258
GENERAL INFORMATION:
APPLICANT: Kwiatkowski, David J.
APPLICANT: Sampson, Julian R.
APPLICANT: Povey, Sue
APPLICANT: van Slegtenhorst, Marjon
APPLICANT: Halley, Dicky
TITLE OF INVENTION: Compositions and Methods Based Upon the Tubercous
TITLE OF INVENTION: Sclerosis-1 (TSC-1) Gene and Gene Product
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSER: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Releasee #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/950,046A
FILING DATE: 12-Sep-2001
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/42002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6585
TELEFAX: (202) 639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1164 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-950-046A-2

Query Match
Best Local Similarity 73.5%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIILLC 9
Db 253 HDVIVLC 259

STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
US-09-457-708-2

Query Match
Best Local Similarity 73.5%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIILLC 9
Db 253 HDVIVLC 259

RESULT 26
US-09-950-046A-2
Sequence 2, Application US/09950046A
Patent No. 6548258
GENERAL INFORMATION:
APPLICANT: Kwiatkowski, David J.
APPLICANT: Sampson, Julian R.
APPLICANT: Povey, Sue
APPLICANT: van Slegtenhorst, Marjon
APPLICANT: Halley, Dicky
TITLE OF INVENTION: Compositions and Methods Based Upon the Tubercous
TITLE OF INVENTION: Sclerosis-1 (TSC-1) Gene and Gene Product
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSER: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Releasee #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/950,046A
FILING DATE: 12-Sep-2001
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/42002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6585
TELEFAX: (202) 639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1164 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-950-046A-2

Query Match
Best Local Similarity 73.5%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIILLC 9
Db 253 HDVIVLC 259

RESULT 27
US-09-976-594-989
; Sequence 989, Application US/0976594
; Patent No. 6673549
; GENERAL INFORMATION:
; APPLICANT: Furness, Michael
; APPLICANT: Buchbinder, Jenny
; TITLE OF INVENTION: GENES EXPRESSED IN C3A LIVER CELL CULTURES TREATED WITH STEROIDS
; FILE REFERENCE: PA-0041 US
; CURRENT APPLICATION NUMBER: US/09/976,594
; CURRENT FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: 60/240,409
; PRIOR FILING DATE: 2000-10-12
; NUMBER OF SEQ ID NOS: 1143
; SOFTWARE: PERL Program
; SEQ ID NO 989
; LENGTH: 1164
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc feature
; OTHER INFORMATION: Incyte ID No. 6673549 4215034CD1
US-09-976-594-989

Query Match 73.5%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 IHDIIEC 9
Db 253 HDVVIC 259

RESULT 28
US-08-934-915-159
; Sequence 159, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1, 5, 6, 8,
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
; TITLE OF INVENTION: DIAGNOSTIC PURPOSES
; NUMBER OF SEQUENCES: 193
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft Word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. FOUTCH
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946.6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:

; INFORMATION FOR SEQ ID NO: 159:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-934-915-159

Query Match 71.4%; Score 35; DB 1; Length 20;
Best Local Similarity 87.5%; Pred. No. 4.1;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TIHDIIE 8
Db 13 TIHDIIE 20

RESULT 29
US-09-270-767-33608
; Sequence 33608, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 33608
; LENGTH: 111
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-33608

Query Match 69.4%; Score 34; DB 2; Length 111;
Best Local Similarity 50.0%; Pred. No. 38;
Matches 4; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IHDIIEC 9
Db 33 LHDILIC 40

RESULT 30
US-09-270-767-48825
; Sequence 48825, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 48825
; LENGTH: 111
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-48825

Query Match 69.4%; Score 34; DB 2; Length 111;
Best Local Similarity 50.0%; Pred. No. 38;
Matches 4; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IHDIIEC 9
Db 33 LHDILIC 40

RESULT 31
US-09-489-039A-13381

Sequence 13381, Application US/09489039A
Patent No. 6610836
GENERAL INFORMATION:
APPLICANT: Gary Breton et. al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
FILE REFERENCE: 2709.2004001
CURRENT APPLICATION NUMBER: US/09/489,039A
CURRENT FILING DATE: 2000-01-27
PRIOR APPLICATION NUMBER: US 60/117,747
PRIOR FILING DATE: 1999-01-29
NUMBER OF SEQ ID NOS: 14342
SEQ ID NO 13381
LENGTH: 266
TYPE: PRT
ORGANISM: Klebsiella pneumoniae
US-09-489-039A-13381

Query Match 69.4%; Score 34; DB 2; Length 266;
Best Local Similarity 57.1%; Pred. No. 95;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 3 HDIILEC 9
|||:|:|
Db 138 HDLVLDLC 144

RESULT 32
US-09-638-937-2
Sequence 2, Application US/09638937
Patent No. 6593514
GENERAL INFORMATION:
APPLICANT: Cahoon, Edgar B
APPLICANT: Hitz, William D
TITLE OF INVENTION: METHOD FOR THE PRODUCTION OF CALENDIC ACID, AN UNUSUAL
TITLE OF INVENTION: PATTY ACID CONTAINING DELTA-9,10,12 CONJUGATED DOUBLE
BONDS
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/638,937
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: BB-1371-P1
PRIOR FILING DATE: 1999-08-16
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Microsoft Office 97
SEQ ID NO 2
LENGTH: 374
TYPE: PRT
ORGANISM: Calendula officinalis
US-09-638-937-2

Query Match 67.3%; Score 33; DB 2; Length 374;
Best Local Similarity 62.5%; Pred. No. 2.1e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 HDIILEC 9
|||:|:|
Db 47 LHDIIIVTC 54

RESULT 33
US-08-426-509A-12
Sequence 12, Application US/08426509A
Patent No. 6326469
GENERAL INFORMATION:
APPLICANT: Ulrich, Axel
APPLICANT: Gishitzky, Mikhail
APPLICANT: Sures, Itman G.
TITLE OF INVENTION: NOVEL MEGAKARYOCYTIC PROTEIN
TITLE OF INVENTION: TYROSINE KINASES
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Pennie & Edmonds

STREET: 1155 Avenue of the Americas
CITY: New York,
STATE: NY
COUNTRY: USA
ZIP: 10036-2711
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/426,509A
FILING DATE: 21-APR-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/232,545
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7683-0074-999
TELECOMMUNICATION INFORMATION:
TELEPHONE: 212-790-9090
TELEFAX: 212-869-9741
TELEX: 66141 PENNIR
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 536 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: No. 6326469e
US-08-426-509A-12

Query Match 67.3%; Score 33; DB 2; Length 536;
Best Local Similarity 33.3%; Pred. No. 3e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

QY 1 THDIILEC 9
|||:|:|
Db 493 SLHDVWVC 501

RESULT 34
US-08-232-545-12
Sequence 12, Application US/08232545
Patent No. 6506578
GENERAL INFORMATION:
APPLICANT: Ulrich, Axel
APPLICANT: Gishitzky, Mikhail
APPLICANT: Sures, Itman G.
TITLE OF INVENTION: No. 6506578e1 Megakaryocytic Protein Tyrosine
TITLE OF INVENTION: Kinases
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York
STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/232,545
FILING DATE: 22-APR-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742

REFERENCE/DOCKET NUMBER: 7683-050
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)790-9090
TELEFAX: (212)869-9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 536 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
US-08-232-545-12

Query Match 67.3%; Score 33; DB 2; Length 536;
Best Local Similarity 33.3%; Pred. No. 3e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TTHDILLEC 9
Db 493 SLHDVMVQC 501

RESULT 35
US-09-977-261-12
Sequence 12, Application US/09977261
Patent No. 6908984
GENERAL INFORMATION:
APPLICANT: ULIRICH, AXEL
APPLICANT: GISHIZKY, MIKHAIL
APPLICANT: SURES, IRMINGARD
TITLE OF INVENTION: NOVEL MEGAKARYOCYTIC PROTEIN TYROSINE KINASES
FILE REFERENCE: 038602/1259
CURRENT APPLICATION NUMBER: US/09/977,261
CURRENT FILING DATE: 2001-10-16
PRIOR APPLICATION NUMBER: 08/232,545
PRIOR FILING DATE: 1994-04-22
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 12
LENGTH: 536
TYPE: PRT
ORGANISM: Gallus gallus
US-09-977-261-12

Query Match 67.3%; Score 33; DB 2; Length 536;
Best Local Similarity 33.3%; Pred. No. 3e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TTHDILLEC 9
Db 493 SLHDVMVQC 501

RESULT 36
PCT-US95-05008-12
Sequence 12, Application PC/TUS9505008
GENERAL INFORMATION:
APPLICANT: Sugan, Inc.
APPLICANT: 515 Galveston Drive
APPLICANT: Redwood City, California 94063-4720
APPLICANT: United States of America
APPLICANT: Wissenschaften E.V.
APPLICANT: Hofgarten Str. 2
APPLICANT: Munchen 80539
APPLICANT: Germany
TITLE OF INVENTION: Novel Megakaryocytic Protein Tyrosine
TITLE OF INVENTION: Kinases
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITY: New York

STATE: New York
COUNTRY: U.S.A.
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/05008
FILING DATE: 24-APR-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/232,545
FILING DATE: 22-APR-1994
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Coruzzi, Laura A.
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 7683-074
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212)790-9090
TELEFAX: (212)869-9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 536 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: protein
PCT-US95-05008-12

Query Match 67.3%; Score 33; DB 4; Length 536;
Best Local Similarity 33.3%; Pred. No. 3e+02;
Matches 3; Conservative 6; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TTHDILLEC 9
Db 493 SLHDVMVQC 501

RESULT 37
US-08-159-339A-573
Sequence 573, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esteban
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESSES:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992

APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 573:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-573

Query Match 65.3%; Score 32; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 7.4;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 DIIIEC 9
Db 1 DIIIEC 6

RESULT 38
US-09-270-767-60201
Sequence 60201, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 60201
LENGTH: 217
TYPE: PRT
ORGANISM: Drosophila melanogaster
FEATURE:
OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-60201

Query Match 65.3%; Score 32; DB 2; Length 217;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 2 IIDIIIEC 9
Db 52 IIDIIIEC 59

RESULT 39
US-09-252-991A-29181
Sequence 29181, Application US/09252991A
Patent No. 6551795
GENERAL INFORMATION:
APPLICANT: Marc J. Rubenfield et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
FILE REFERENCE: 107196.136
CURRENT APPLICATION NUMBER: US/09/252,991A
CURRENT FILING DATE: 1999-02-18
PRIORITY FILING DATE: 1998-02-18
PRIORITY FILING DATE: 1998-02-18
PRIORITY APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27
NUMBER OF SEQ ID NOS: 33142
SEQ ID NO 29181
LENGTH: 227
TYPE: PRT
ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-29181

Query Match 65.3%; Score 32; DB 2; Length 227;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 2 IIDIIIEC 9
Db 140 IIDFVNEC 147

RESULT 40
US-09-107-532A-5486
Sequence 5486, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
NUMBER OF SEQUENCES: 7310
CORRESPONDENCE ADDRESS:
ADDRESS: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
CITY: Waltham
STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD-ROM ISO9660
COMPUTER: PC
OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/107,532A
FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/085,598
FILING DATE: 14 May 1998
APPLICATION NUMBER: 60/051571
FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ariniello, Pamela Deneke
REGISTRATION NUMBER: 40,489
REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
TELEPHONE: (781) 893-5007
TELEFAX: (781) 893-8277
INFORMATION FOR SEQ ID NO: 5486:
SEQUENCE CHARACTERISTICS:
LENGTH: 305 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
ORGANISM: Enterococcus faecium
FEATURE:
NAME/KEY: misc feature
LOCATION: (B) LOCATION 1...305
SEQUENCE DESCRIPTION: SEQ ID NO: 5486:
US-09-107-532A-5486

Query Match 65.3%; Score 32; DB 2; Length 305;
Best Local Similarity 100.0%; Pred. No. 2.6e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 4 DIIIEC 9

Db 129 DIIIEC 134

RESULT 41

US-09-266-965-112
; Sequence 112, Application US/09266965
; Patent No. 6495348
; GENERAL INFORMATION:
; APPLICANT: Sherman, D
; APPLICANT: Mao, Y
; APPLICANT: Varoglu, M
; APPLICANT: He, M
; APPLICANT: Sheldon, P
; TITLE OF INVENTION: Mitomycin biosynthetic gene cluster
; FILE REFERENCE: 600.456US1
; CURRENT APPLICATION NUMBER: US/09/266,965
; CURRENT FILING DATE: 1999-03-12
; EARLIER APPLICATION NUMBER: US 08/624,447
; EARLIER FILING DATE: 1996-08-19
; EARLIER APPLICATION NUMBER: PCT/US94/11279
; EARLIER FILING DATE: 1994-10-06
; EARLIER APPLICATION NUMBER: US 08/133,963
; EARLIER FILING DATE: 1993-10-07
; NUMBER OF SEQ ID NOS: 145
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 112
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Streptomyces lavendulae
US-09-266-965-112

Query Match 65.3%; Score 32; DB 2; Length 343;

Best Local Similarity 44.4%; Pred. No. 2.9e+02; Mismatches 2; Indels 0; Gaps 0;

Qy 1 TIHDIIEC 9
Db 90 TRHDVVC 98

RESULT 42

US-09-248-796A-22434
; Sequence 22434, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstein et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 22434
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-22434

Query Match 65.3%; Score 32; DB 2; Length 343;

Best Local Similarity 100.0%; Pred. No. 2.9e+02; Mismatches 0; Indels 0; Gaps 0;

Qy 4 DIIIEC 9
Db 122 DIIIEC 127

RESULT 43

US-09-270-767-44745
; Sequence 44745, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 44745
; LENGTH: 361
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-44745

Query Match 65.3%; Score 32; DB 2; Length 361;

Best Local Similarity 75.0%; Pred. No. 3.1e+02; Mismatches 2; Indels 0; Gaps 0;

Qy 2 IHDIIEC 9
Db 196 IHTIIEC 203

RESULT 44

US-09-248-796A-15186
; Sequence 15186, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstein et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 15186
; LENGTH: 512
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-15186

Query Match 65.3%; Score 32; DB 2; Length 512;

Best Local Similarity 33.3%; Pred. No. 4.5e+02; Mismatches 1; Indels 0; Gaps 0;

Qy 1 TIHDIIEC 9
Db 215 TLVDVWNC 223

RESULT 45

US-09-527-073-2
; Sequence 2, Application US/09527073
; Patent No. 6534313
; GENERAL INFORMATION:
; APPLICANT: Michael M. Neff
; APPLICANT: Joanne Cho Y
; TITLE OF INVENTION: GENETICALLY MODIFIED PLANTS HAVING
; FILE REFERENCE: SALKINS 024A
; CURRENT APPLICATION NUMBER: US/09/527,073
; CURRENT FILING DATE: 2000-03-16
; PRIOR APPLICATION NUMBER: US 60/124570
; PRIOR FILING DATE: 1999-03-16

PRIOR APPLICATION NUMBER: US 60/170,931
PRIOR FILING DATE: 1999-12-14
PRIOR APPLICATION NUMBER: US 60/172,832
PRIOR FILING DATE: 1999-12-20
NUMBER OF SEQ ID NOS: 16
SOFTWARE: fastseq for windows version 4.0
SEQ ID NO 2
LENGTH: 520
TYPE: PRT
ORGANISM: Arabidopsis thaliana
US-09-527-073-2

Query Match 65.3%; Score 32; DB 2; Length 520;
Best Local Similarity 55.6%; Pred. No. 4.5e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 THDIIIEC 9
Db 310 TVQDIVERC 318

RESULT 46
US-08-448-196A-9
Sequence 9, Application US/08448196A
Patent No. 5780594
GENERAL INFORMATION:
APPLICANT: CARTER, DANIEL C.
TITLE OF INVENTION: BIOLOGICALLY ACTIVE PROTEIN FRAGMENTS
CONTAINING SPECIFIC BINDING REGIONS OF SERUM ALBUMIN OR
TITLE OF INVENTION: RELATED PROTEINS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: NASA
STREET: MARSHALL SPACE FLIGHT CENTER
CITY: HUNTSVILLE
STATE: ALABAMA
COUNTRY: USA
ZIP: 35812
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/448,196A
FILING DATE: 23-MAY-1995
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: BROAD JR., ROBERT L.
REGISTRATION NUMBER: 18,757
REFERENCE/DOCKET NUMBER: XX/MFS-28402-2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 205-544-0021
TELEFAX: 205-544-0258
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 590 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULAR TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
US-08-448-196A-9

Query Match 65.3%; Score 32; DB 1; Length 590;
Best Local Similarity 44.4%; Pred. No. 5.2e+02;
Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

QY 1 THDIIIEC 9
Db 433 TVHDIILAC 441

RESULT 47
US-09-785-381-11
Sequence 11, Application US/09785381
Patent No. 6602992
GENERAL INFORMATION:
APPLICANT: DALLOS, Peter
APPLICANT: ZHENG, Jing
APPLICANT: MADISON, Laird
TITLE OF INVENTION: A MAMMALIAN PRESTIN
FILE REFERENCE: 0290-37U1
CURRENT APPLICATION NUMBER: US/09/785,381
PRIOR FILING DATE: 2001-02-15
PRIOR APPLICATION NUMBER: US 60/183,461
PRIOR FILING DATE: 2000-02-18
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn version 3.0
SEQ ID NO 11
LENGTH: 780
TYPE: PRT
ORGANISM: Rattus norvegicus
US-09-785-381-11

Query Match 65.3%; Score 32; DB 2; Length 780;
Best Local Similarity 50.0%; Pred. No. 6.9e+02;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 2 IHDIILIEC 9
Db 655 IHSVLVDC 662

RESULT 48
US-09-949-016-11498
Sequence 11498, Application US/09949016
Patent No. 6812339
GENERAL INFORMATION:
APPLICANT: VENTER, J. Craig et al.
TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
FILE REFERENCE: CL001307
CURRENT APPLICATION NUMBER: US/09/949,016
CURRENT FILING DATE: 2000-04-14
PRIOR APPLICATION NUMBER: 60/241,755
PRIOR FILING DATE: 2000-10-20
PRIOR APPLICATION NUMBER: 60/237,768
PRIOR FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 60/231,498
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 207012
SOFTWARE: fastseq for windows version 4.0
SEQ ID NO 11498
LENGTH: 803
TYPE: PRT
ORGANISM: Human
US-09-949-016-11498

Query Match 65.3%; Score 32; DB 2; Length 803;
Best Local Similarity 50.0%; Pred. No. 7.1e+02;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 2 IHDIILIEC 9
Db 678 IHSVLVDC 685

RESULT 49
US-09-248-796A-15476
Sequence 15476, Application US/09248796A
Patent No. 6747137
GENERAL INFORMATION:
APPLICANT: Keith Weinstock et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICANS

TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 15476
LENGTH: 1060
TYPE: PRT
ORGANISM: Candida albicans
US-09-248-796A-15476

Query Match 65.3%; Score 32; DB 2; Length 1060;
Best Local Similarity 75.0%; Pred. No. 9.5e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 TTHDILE 8
Db 49 TTHDLSLE 56

RESULT 50
US-08-787-547-102
Sequence 102; Application US/08787547
Patent No. 5783567
GENERAL INFORMATION:
APPLICANT: Hedley, Mary Lynne
APPLICANT: Curley, Joanne M.
APPLICANT: Langer, Robert S.
TITLE OF INVENTION: MICROPARTICLES FOR DELIVERY
TITLE OF INVENTION: OF NUCLEIC ACID
NUMBER OF SEQUENCES: 107
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
CITY: Boston
STATE: MA
COUNTRY: US
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/787,547
FILING DATE: 22-JAN-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Praeger, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 08191/003001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-5070
TELEFAX: 617-542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 102:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-787-547-102

Query Match 63.3%; Score 31; DB 1; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDII 6
Db 4 TTHDII 9

Search completed: May 5, 2006, 04:01:02
Job time : 24 secs

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:07:45 ; Search time 55.8 Seconds
(without alignments)
67.392 Million cell updates/sec

Title: US-08-170-344-7
Perfect score: 49
Sequence: 1 T1HD1LEC 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Published Applications AA Main:*

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2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	ID	Description
1	49	100.0	9	4	US-10-777-053-547 Sequence 547, App
2	49	100.0	9	4	US-10-837-217-547 Sequence 547, App
3	49	100.0	10	4	US-10-128-711-72 Sequence 72, App1
4	49	100.0	10	4	US-10-133-210-280 Sequence 280, App
5	49	100.0	15	4	US-10-476-570-22 Sequence 22, App1
6	49	100.0	15	4	US-10-476-570-22 Sequence 23, App1
7	49	100.0	30	4	US-10-476-570-53 Sequence 53, App1
8	49	100.0	30	5	US-10-858-384-4 Sequence 4, App1
9	49	100.0	32	4	US-10-476-570-9 Sequence 9, App1
10	49	100.0	33	4	US-10-476-570-19 Sequence 19, App1
11	49	100.0	151	5	US-10-177-390-6 Sequence 6, App1
12	49	100.0	151	5	US-10-484-063-20 Sequence 20, App1
13	49	100.0	151	5	US-10-884-063-27 Sequence 27, App1
14	49	100.0	158	5	US-10-858-384-2 Sequence 2, App1
15	49	100.0	158	5	US-10-367-057-16 Sequence 16, App1
16	49	100.0	158	6	US-11-021-949-13 Sequence 13, App1
17	49	100.0	171	4	US-10-472-724-2 Sequence 2, App1
18	49	100.0	263	6	US-11-072-288-1 Sequence 1, App1
19	49	100.0	266	3	US-09-367-309A-1 Sequence 1, App1
20	49	100.0	273	4	US-10-000-903-4 Sequence 4, App1
21	49	100.0	273	5	US-10-899-771-4 Sequence 4, App1
22	49	100.0	292	4	US-10-000-903-10 Sequence 10, App1
23	49	100.0	292	5	US-10-899-771-10 Sequence 10, App1
24	49	100.0	371	4	US-10-000-903-6 Sequence 6, App1
25	49	100.0	371	5	US-10-899-771-6 Sequence 6, App1
26	49	100.0	390	4	US-10-000-903-14 Sequence 14, App1
27	49	100.0	390	5	US-10-899-771-14 Sequence 14, App1

28	49	100.0	536	4	US-10-367-095-10 Sequence 10, App1
29	49	100.0	536	4	US-10-368-046-10 Sequence 10, App1
30	49	100.0	536	4	US-10-367-367-10 Sequence 10, App1
31	49	100.0	536	5	US-10-918-337-10 Sequence 10, App1
32	44	89.8	21	4	US-10-476-570-10 Sequence 10, App1
33	41	83.7	306	4	US-10-437-963-149433 Sequence 24, App1
34	40	81.6	15	4	US-10-476-570-24 Sequence 24, App1
35	40	81.6	150	4	US-10-437-963-153797 Sequence 153797, App1
36	39	79.6	297	4	US-10-424-589-275411 Sequence 165507, App1
37	39	79.6	298	4	US-10-424-589-168507 Sequence 21847, App1
38	37	75.5	145	5	US-10-732-923-21847 Sequence 361, App
39	37	75.5	158	6	US-11-021-949-361 Sequence 10846, App
40	37	75.5	554	5	US-10-739-930-10846 Sequence 33593, App
41	37	75.5	70	4	US-10-029-386-33593 Sequence 205547, App
42	36	73.5	83	4	US-10-425-115-205547 Sequence 194068, App
43	36	73.5	117	4	US-10-425-115-194068 Sequence 31, App1
44	36	73.5	162	6	US-11-021-949-31 Sequence 358720, App
45	36	73.5	164	4	US-10-425-115-358720 Sequence 358724, App1
46	36	73.5	173	4	US-10-425-115-358724 Sequence 18, App1
47	36	73.5	174	5	US-10-947-979-18 Sequence 236213, App1
48	36	73.5	176	4	US-10-425-115-236213 Sequence 10, App1
49	36	73.5	211	4	US-10-947-979-10 Sequence 56403, App
50	36	73.5	216	4	US-10-425-114-56403 Sequence 45686, App
51	36	73.5	220	4	US-10-767-701-45686 Sequence 58813, App
52	36	73.5	274	4	US-10-425-114-58813 Sequence 65774, App
53	36	73.5	274	4	US-10-425-114-65774 Sequence 145297, App
54	36	73.5	291	4	US-10-437-963-145297 Sequence 52907, App
55	36	73.5	294	4	US-10-425-114-52907 Sequence 345578, App
56	36	73.5	297	4	US-10-767-701-36891 Sequence 36891, App
57	36	73.5	304	4	US-10-425-115-345578 Sequence 126120, App
58	36	73.5	306	4	US-10-437-963-126120 Sequence 50748, App
59	36	73.5	312	4	US-10-425-114-50748 Sequence 151133, App
60	36	73.5	356	4	US-10-437-963-151133 Sequence 159590, App
61	36	73.5	403	4	US-10-437-963-195950 Sequence 195950, App
62	36	73.5	501	4	US-10-437-963-153537 Sequence 96, App1
63	36	73.5	789	4	US-10-342-844-96 Sequence 13152, App
64	36	73.5	1100	6	US-11-097-143-1152 Sequence 2, App1
65	36	73.5	1164	3	US-09-950-046A-2 Sequence 793, App
66	36	73.5	1164	3	US-10-719-993-793 Sequence 794, App
67	36	73.5	1164	5	US-10-719-993-794 Sequence 795, App
68	36	73.5	1164	5	US-10-719-993-795 Sequence 3779, App
69	36	73.5	2539	4	US-10-369-493-3779 Sequence 261856, App
70	35	71.4	76	4	US-10-424-599-261856 Sequence 135239, App
71	35	71.4	108	4	US-10-437-963-135239 Sequence 19, App1
72	35	71.4	148	6	US-11-021-949-19 Sequence 3636, App
73	35	71.4	180	6	US-11-097-143-36396 Sequence 121171, App
74	35	71.4	182	4	US-10-437-963-121171 Sequence 39716, App
75	35	71.4	271	4	US-10-767-701-39716 Sequence 187372, App
76	35	71.4	294	4	US-10-424-599-187372 Sequence 193535, App
77	35	71.4	428	4	US-10-425-115-193535 Sequence 106, App
78	35	71.4	527	4	US-10-078-923-106 Sequence 54439, App
79	35	71.4	548	4	US-10-425-115-54439 Sequence 278342, App
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81	35	71.4	570	4	US-10-437-963-116928 Sequence 201770, App
82	35	71.4	788	4	US-10-437-963-201770 Sequence 237714, App
83	35	71.4	1025	4	US-10-425-115-237714 Sequence 344428, App
84	35	71.4	1047	4	US-10-425-115-344428 Sequence 256316, App
85	35	71.4	1145	4	US-10-425-115-256316 Sequence 239507, App
86	35	71.4	1291	4	US-10-425-115-239207 Sequence 344339, App
87	35	71.4	1388	4	US-10-425-115-344339 Sequence 344399, App
88	35	71.4	1794	4	US-10-425-115-344399 Sequence 239216, App
89	35	71.4	1848	4	US-10-425-115-239216 Sequence 344401, App
90	35	71.4	1869	4	US-10-425-115-344401 Sequence 359998, App
91	34	69.4	107	4	US-10-425-115-359998 Sequence 62097, App
92	34	69.4	110	4	US-10-767-701-62097 Sequence 226149, App
93	34	69.4	117	4	US-10-425-115-226149 Sequence 6393, App
94	34	69.4	123	6	US-11-097-143-6390 Sequence 274871, App
95	34	69.4	148	4	US-10-425-115-274871 Sequence 16, App1
96	34	69.4	149	6	US-11-021-949-16 Sequence 1123, App
97	34	69.4	155	3	US-09-925-299-1123 Sequence 43474, App
98	34	69.4	165	4	US-10-767-701-43474 Sequence 270, App
99	34	69.4	217	4	US-10-474-776-270 Sequence 270, App
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258	32	65.3	743	4	US-10-287-226-358	Sequence 368, App	331	31	63.3	267	3	US-09-981-528-2	Sequence 14, Appl
259	32	65.3	774	4	US-10-437-963-115395	Sequence 115395, A	332	31	63.3	274	3	US-09-790-045-14	Sequence 14, Appl
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262	32	65.3	780	5	US-10-631-467-14	Sequence 14, Appl	335	31	63.3	275	3	US-10-021-660-98	Sequence 3, Appl
263	32	65.3	780	5	US-10-631-467-667	Sequence 667, App	336	31	63.3	275	4	US-10-021-660-98	Sequence 64, Appl
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269	32	65.3	1071	4	US-10-437-963-176644	Sequence 176644, A	342	31	63.3	275	5	US-10-883-576-59	Sequence 1942, Ap
270	32	65.3	1264	4	US-10-032-588-7118	Sequence 7118, Ap	343	31	63.3	279	4	US-10-369-493-1942	Sequence 274112, A
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298	31	63.3					371	31	63.3	384	4	US-10-425-114-65198	Sequence 53294, A
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442	31	63.3	4097	4	US-10-408-765A-1598	Sequence 1598, Ap
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444	31	63.3	4128	4	US-10-363-616-416	Sequence 416, App
445	31	63.3	4128	4	US-10-764-425-152	Sequence 152, App
446	31	63.3	4128	5	US-10-473-127-568	Sequence 568, App
447	31	63.3	4128	5	US-10-473-127-573	Sequence 573, App
448	31	63.3	4128	5	US-10-473-127-574	Sequence 574, App
449	31	63.3	4128	5	US-10-511-561-3	Sequence 3, Appl1
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452	31	63.3	5175	4	US-10-369-493-6861	Sequence 6861, Ap
453	31	63.3	5198	4	US-10-120-801-75	Sequence 75, Appl
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455	31	63.3	5198	4	US-10-369-493-6860	Sequence 6860, Ap
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508	30	61.2	205	4	US-10-369-493-966	Sequence 966, App
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512	30	61.2	205	5	US-10-486-645-5	Sequence 5, Appl1
513	30	61.2	205	5	US-10-929-759-243	Sequence 243, App
514	30	61.2	205	5	US-10-973-919-243	Sequence 243, App
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533	30	61.2	257	4	US-10-218-779-100	Sequence 100, App
534	30	61.2	258	3	US-09-964-956-75	Sequence 75, Appl
535	30	61.2	258	4	US-10-042-865-100	Sequence 100, App
536	30	61.2	258	4	US-10-072-012-801	Sequence 801, App
537	30	61.2	258	4	US-10-072-012-858	Sequence 858, App
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632	30	61.2	522	4	US-10-138-916-96	Sequence 96, Appl
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634	30	61.2	522	4	US-10-139-218-96	Sequence 96, Appl
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636	30	61.2	522	4	US-10-138-898-96	Sequence 96, Appl
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646	30	61.2	526	5	US-10-732-923-13493	Sequence 13493, A
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648	30	61.2	526	5	US-10-732-923-13499	Sequence 13499, A
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652	30	61.2	533	4	US-10-321-802-8	Sequence 8, Appl
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658	30	61.2	535	4	US-10-424-599-210382	Sequence 210382, A
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662	30	61.2	536	3	US-09-929-266-10	Sequence 10, Appl
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664	30	61.2	536	3	US-09-977-261-13	Sequence 2, Appl
665	30	61.2	536	3	US-10-691-079-2	Sequence 2, Appl
666	30	61.2	536	5	US-10-887-588-2	Sequence 2, Appl
667	30	61.2	536	5	US-10-497-641-2	Sequence 2, Appl
668	30	61.2	536	5	US-10-825-566-10	Sequence 10, Appl
669	30	61.2	537	4	US-10-321-802-32	Sequence 32, Appl
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689	30	61.2	590	5	US-10-484-063-21	Sequence 21, Appl	762	30	61.2	945	4	US-10-131-837A-146	Sequence 146, App
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693	30	61.2	635	3	US-09-815-242-11209	Sequence 11209, A	766	30	61.2	945	4	US-10-147-515-146	Sequence 146, App
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695	30	61.2	636	4	US-10-104-047-2449	Sequence 2449, Ap	768	30	61.2	945	4	US-10-147-526-146	Sequence 146, App
696	-30	61.2	667	5	US-10-450-763-52279	Sequence 52279, A	769	30	61.2	945	4	US-10-147-527-146	Sequence 146, App
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698	30	61.2	709	4	US-10-437-963-116285	Sequence 116285,	771	30	61.2	945	4	US-10-121-043-146	Sequence 146, App
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702	-30	61.2	772	5	US-10-733-923-13485	Sequence 13485, A	775	30	61.2	945	4	US-10-123-908-146	Sequence 146, App
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705	30	61.2	837	4	US-10-282-132A-77952	Sequence 77952, A	778	30	61.2	945	4	US-10-124-813-146	Sequence 146, App
706	30	61.2	857	4	US-10-437-963-173301	Sequence 173301,	779	30	61.2	945	4	US-10-124-817-146	Sequence 146, App
707	30	61.2	906	4	US-10-437-963-128708	Sequence 128708,	780	30	61.2	945	4	US-10-125-922-146	Sequence 146, App
708	30	61.2	912	4	US-10-437-963-177936	Sequence 177936,	781	30	61.2	945	4	US-10-125-924-146	Sequence 146, App
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752	30	61.2	945	4	US-10-127-829A-146	Sequence 146, App	825	30	61.2	945	4	US-10-125-928A-146	Sequence 146, App
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851	30	61.2	945	4	US-10-137-865A-146	Sequence 146, App	924	30	61.2	945	4	US-10-152-531-146	Sequence 146, App
852	30	61.2	945	4	US-10-137-866A-146	Sequence 146, App	925	30	61.2	945	4	US-10-127-840A-146	Sequence 146, App
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856	30	61.2	945	4	US-10-121-042-146	Sequence 146, App	929	30	61.2	945	4	US-10-142-887-146	Sequence 146, App
857	30	61.2	945	4	US-10-123-912-146	Sequence 146, App	930	30	61.2	945	4	US-10-142-888-146	Sequence 146, App
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864	30	61.2	945	4	US-10-146-728-146	Sequence 146, App	937	30	61.2	945	4	US-10-145-091-146	Sequence 146, App
865	30	61.2	945	4	US-10-152-380-146	Sequence 146, App	938	30	61.2	945	4	US-10-145-629-146	Sequence 146, App
866	30	61.2	945	4	US-10-153-934-146	Sequence 146, App	939	30	61.2	945	4	US-10-145-630-146	Sequence 146, App
867	30	61.2	945	4	US-10-140-807-146	Sequence 146, App	940	30	61.2	945	4	US-10-145-747-146	Sequence 146, App
868	30	61.2	945	4	US-10-140-924-146	Sequence 146, App	941	30	61.2	945	4	US-10-145-752-146	Sequence 146, App
869	30	61.2	945	4	US-10-140-926-146	Sequence 146, App	942	30	61.2	945	4	US-10-145-755-146	Sequence 146, App
870	30	61.2	945	4	US-10-141-638-146	Sequence 146, App	943	30	61.2	945	4	US-10-145-818-146	Sequence 146, App
871	30	61.2	945	4	US-10-141-702-146	Sequence 146, App	944	30	61.2	945	4	US-10-145-820-146	Sequence 146, App
872	30	61.2	945	4	US-10-141-704-146	Sequence 146, App	945	30	61.2	945	4	US-10-145-824-146	Sequence 146, App
873	30	61.2	945	4	US-10-142-421-146	Sequence 146, App	946	30	61.2	945	4	US-10-145-827-146	Sequence 146, App
874	30	61.2	945	4	US-10-142-432-146	Sequence 146, App	947	30	61.2	945	4	US-10-147-481-146	Sequence 146, App
875	30	61.2	945	4	US-10-142-757-146	Sequence 146, App	948	30	61.2	945	4	US-10-147-482-146	Sequence 146, App
876	30	61.2	945	4	US-10-143-033-146	Sequence 146, App	949	30	61.2	945	4	US-10-147-503-146	Sequence 146, App
877	30	61.2	945	4	US-10-144-994-146	Sequence 146, App	950	30	61.2	945	4	US-10-147-522-146	Sequence 146, App
878	30	61.2	945	4	US-10-145-628-146	Sequence 146, App	951	30	61.2	945	4	US-10-152-401-146	Sequence 146, App
879	30	61.2	945	4	US-10-145-746-146	Sequence 146, App	952	30	61.2	945	4	US-10-157-783-146	Sequence 146, App
880	30	61.2	945	4	US-10-145-748-146	Sequence 146, App	953	30	61.2	945	4	US-10-158-792-146	Sequence 146, App
881	30	61.2	945	4	US-10-145-823-146	Sequence 146, App	954	30	61.2	945	4	US-10-145-824-146	Sequence 146, App
882	30	61.2	945	4	US-10-145-826-146	Sequence 146, App	955	30	61.2	945	4	US-10-145-827-146	Sequence 146, App
883	30	61.2	945	4	US-10-145-870-146	Sequence 146, App	956	30	61.2	945	4	US-10-145-829-146	Sequence 146, App
884	30	61.2	945	4	US-10-145-876-146	Sequence 146, App	957	30	61.2	945	4	US-10-145-831-146	Sequence 146, App
885	30	61.2	945	4	US-10-145-959-146	Sequence 146, App	958	30	61.2	945	4	US-10-145-835-146	Sequence 146, App
886	30	61.2	945	4	US-10-146-724-146	Sequence 146, App	959	30	61.2	945	4	US-10-145-837-146	Sequence 146, App
887	30	61.2	945	4	US-10-146-725-146	Sequence 146, App	960	30	61.2	945	4	US-10-146-790-146	Sequence 146, App
888	30	61.2	945	4	US-10-146-795-146	Sequence 146, App	961	30	61.2	945	4	US-10-146-793-146	Sequence 146, App
889	30	61.2	945	4	US-10-147-495-146	Sequence 146, App	962	30	61.2	945	4	US-10-147-480-146	Sequence 146, App
890	30	61.2	945	4	US-10-147-501-146	Sequence 146, App	963	30	61.2	945	4	US-10-147-486-146	Sequence 146, App
891	30	61.2	945	4	US-10-147-504-146	Sequence 146, App	964	30	61.2	945	4	US-10-147-487-146	Sequence 146, App
892	30	61.2	945	4	US-10-147-506-146	Sequence 146, App	965	30	61.2	945	4	US-10-147-490-146	Sequence 146, App
893	30	61.2	945	4	US-10-147-509-146	Sequence 146, App	966	30	61.2	945	4	US-10-147-494-146	Sequence 146, App
894	30	61.2	945	4	US-10-147-510-146	Sequence 146, App	967	30	61.2	945	4	US-10-147-498-146	Sequence 146, App
895	30	61.2	945	4	US-10-147-511-146	Sequence 146, App	968	30	61.2	945	4	US-10-147-514-146	Sequence 146, App
896	30	61.2	945	4	US-10-147-529-146	Sequence 146, App	969	30	61.2	945	4	US-10-147-485-146	Sequence 146, App
897	30	61.2	945	4	US-10-152-397-146	Sequence 146, App	970	30	61.2	945	4	US-10-147-486-146	Sequence 146, App
898	30	61.2	945	4	US-10-153-586-146	Sequence 146, App	971	30	61.2	945	4	US-10-147-487-146	Sequence 146, App
899	30	61.2	945	4	US-10-158-786-146	Sequence 146, App	972	30	61.2	945	4	US-10-147-490-146	Sequence 146, App
900	30	61.2	945	4	US-10-137-870-146	Sequence 146, App	973	30	61.2	945	4	US-10-147-494-146	Sequence 146, App
901	30	61.2	945	4	US-10-140-018-146	Sequence 146, App	974	30	61.2	945	4	US-10-147-498-146	Sequence 146, App
902	30	61.2	945	4	US-10-140-021-146	Sequence 146, App	975	30	61.2	945	4	US-10-147-514-146	Sequence 146, App
903	30	61.2	945	4	US-10-140-027-146	Sequence 146, App	976	30	61.2	945	4	US-10-147-514-146	Sequence 146, App

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977 30 61.2 945 4 US-10-147-524-146 Sequence 146, App
978 30 61.2 945 4 US-10-152-379-146 Sequence 146, App
979 30 61.2 945 4 US-10-152-394-146 Sequence 146, App
980 30 61.2 945 4 US-10-152-406-146 Sequence 146, App
981 30 61.2 945 4 US-10-156-847-146 Sequence 146, App
982 30 61.2 945 4 US-10-157-778-146 Sequence 146, App
983 30 61.2 945 4 US-10-157-799-146 Sequence 146, App
984 30 61.2 945 4 US-10-160-504-146 Sequence 146, App
985 30 61.2 945 4 US-10-145-634-146 Sequence 146, App
986 30 61.2 945 4 US-10-147-520-146 Sequence 146, App
987 30 61.2 945 4 US-10-157-781-146 Sequence 146, App
988 30 61.2 945 4 US-10-176-989-146 Sequence 146, App
989 30 61.2 945 4 US-10-147-491-146 Sequence 146, App
990 30 61.2 945 4 US-10-152-378-146 Sequence 146, App
991 30 61.2 945 4 US-10-152-382-146 Sequence 146, App
992 30 61.2 945 4 US-10-152-383-146 Sequence 146, App
993 30 61.2 945 4 US-10-152-384-146 Sequence 146, App
994 30 61.2 945 4 US-10-152-387-146 Sequence 146, App
995 30 61.2 945 4 US-10-152-389-146 Sequence 146, App
996 30 61.2 945 4 US-10-152-390-146 Sequence 146, App
997 30 61.2 945 4 US-10-152-392-146 Sequence 146, App
998 30 61.2 945 4 US-10-153-756-146 Sequence 146, App
999 30 61.2 945 4 US-10-157-779-146 Sequence 146, App
1000 30 61.2 945 4 US-10-157-784-146 Sequence 146, App
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ALIGNMENTS

```
RESULT 1
US-10-777-053-547
; Sequence 547, Application US/10777053
; Publication No. US20040132086A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Diamond, David C.
; APPLICANT: Qiu, Zhiyong
; APPLICANT: Lei, Xiang-Dong
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPTIOPES OF
; FILE REFERENCE: TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
; FILE REFERENCE: MANNA.022C1
; CURRENT FILING DATE: 2004-02-10
; PRIOR APPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: 60/336,968
; NUMBER OF SEQ ID NOS: 979
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 547
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papillomavirus 16
US-10-777-053-547

Query Match 100.0%; Score 49; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPTIOPES OF
; FILE REFERENCE: MANNA.022C2
; CURRENT APPLICATION NUMBER: US/10/837,217
; PRIOR FILING DATE: 2004-04-30
; PRIOR APPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: 60/336,968
; NUMBER OF SEQ ID NOS: 979
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 547
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papillomavirus 16
US-10-837-217-547

Query Match 100.0%; Score 49; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
RESULT 3
US-10-128-711-72
; Sequence 72, Application US/10128711
; Publication No. US2003009634A1
; GENERAL INFORMATION:
; APPLICANT: VITTELLO, Maria A.
; APPLICANT: CHESTNUT, Robert W.
; APPLICANT: SETTE, Alessandro D.
; APPLICANT: CELIS, Esteban
; APPLICANT: GRAY, Howard
; TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
; CTL IMMUNITY
; NUMBER OF SEQUENCES: 153
; CORRESPONDENCE ADDRESS:
; ADDRESSER: Townsend and Townsend Kourile and Crew
; STREET: Stewart Street Tower, One Market Plaza
; CITY: San Francisco
; STATE: California
; COUNTRY: US
; ZIP: 94105-1493
; COMPUTER READABLE FORM:
; MEDIUM TYPE: floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/10/128,711
; FILING DATE: 22-Apr-2002
; CLASSIFICATION: <Unknown>
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US/08/197,484
; FILING DATE: 16-FEB-1994
; APPLICATION NUMBER: US 07/935,811
; FILING DATE: 26-AUG-1992
; APPLICATION NUMBER: US 07/874,491
; FILING DATE: 27-APR-1992
; APPLICATION NUMBER: US 07/827,682
; FILING DATE: 29-JAN-1992
; APPLICATION NUMBER: US 07/749,568
; FILING DATE: 26-AUG-1991
; ATTORNEY/AGENT INFORMATION:
; NAME: Parmelee, Steven W.
; REGISTRATION NUMBER: 31,990
; REFERENCE/DOCKET NUMBER: 14137-26-4
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (206) 467-9600
; TELEFAX: (206) 623-6793
```

CPs

INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 72:
US-10-128-711-72

Query Match 100.0%; Score 49; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
Db 1 TTHDIIIEC 9

RESULT 4
US-10-133-210-280

Sequence 280, Application US/10133210
Publication No. US20030103964A1
GENERAL INFORMATION:
APPLICANT: Delisl, Charles
APPLICANT: Berzofsky, Jay
APPLICANT: Gulukota, Kamalakari
APPLICANT: Vaccaro, Dennis
APPLICANT: Weng, Ziping
APPLICANT: Zhang, Chao
TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
FILE REFERENCE: BU-035AX
CURRENT APPLICATION NUMBER: US/10/133,210
CURRENT FILING DATE: 2002-04-26
NUMBER OF SEQ ID NOS: 281
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 280
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-280

Query Match 100.0%; Score 49; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
Db 1 TTHDIIIEC 9

RESULT 5
US-10-476-570-22

Sequence 22, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLER, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980

PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 22
LENGTH: 15
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 24-38
US-10-476-570-22

Query Match 100.0%; Score 49; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.016;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
Db 6 TTHDIIIEC 14

RESULT 6
US-10-476-570-23

Sequence 23, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLER, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
NUMBER OF SEQ ID NOS: 63
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 23
LENGTH: 15
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 28-42
US-10-476-570-23

Query Match 100.0%; Score 49; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.016;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
Db 2 TTHDIIIEC 10

RESULT 7
US-10-476-570-53

Sequence 53, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLER, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US

```

; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53

Query Match      100.0%; Score 49; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.035;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 THIDIIIEC 9
Db      15 THIDIIIEC 23

RESULT 8
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US2005003025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT-VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCOISE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; CURRENT FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match      100.0%; Score 49; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.035;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 THIDIIIEC 9
Db      15 THIDIIIEC 23

RESULT 9
US-10-476-570-9
; Sequence 9, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: POUVELLE-MORATILLE, Sandra
```

```

; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 9
; LENGTH: 32
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9

Query Match      100.0%; Score 49; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.037;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 THIDIIIEC 9
Db      16 THIDIIIEC 24

RESULT 10
US-10-476-570-19
; Sequence 19, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 19
; LENGTH: 33
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19

Query Match      100.0%; Score 49; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.039;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy      1 THIDIIIEC 9
Db      16 THIDIIIEC 24

RESULT 11
US-10-177-390-6
; Sequence 6, Application US/10177390
; Publication No. US20030143743A1
; GENERAL INFORMATION:
```

APPLICANT: Schuler, Gerold
APPLICANT: N.V. Antwerp Innovatiecentrum
TITLE OF INVENTION: Improved Transfection of Eucaryotic Cells with Linear
TITLE OF INVENTION: Polynucleotides by Electroporation
FILE REFERENCE: 02150500/3H/ml
CURRENT APPLICATION NUMBER: US/10/177,390
CURRENT FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match 100.0%; Score 49; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDIILEC 9
Db 22 T1HDIILEC 30

RESULT 12
US-10-484-063-20
Sequence 20, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLEMO
APPLICANT: FOLLEN, MICHELE
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 20
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-484-063-20

Query Match 100.0%; Score 49; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDIILEC 9
Db 22 T1HDIILEC 30

RESULT 13
US-10-484-063-27
Sequence 27, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLEMO
APPLICANT: FOLLEN, MICHELE
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
TITLE OF INVENTION: PRE-CANCEROUS AND CANCEROUS GROWTHS, INCLUDING CIN
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19

PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-484-063-27

Query Match 100.0%; Score 49; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDIILEC 9
Db 22 T1HDIILEC 30

RESULT 14
US-10-858-384-2
Sequence 2, Application US/10858384
Publication No. US20050033025A1
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCINE
APPLICANT: FERRIER, ESTELLE
TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
TITLE OF INVENTION: PARTICULARLY IN VACCINATION
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858,384
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2

Query Match 100.0%; Score 49; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 T1HDIILEC 9
Db 29 T1HDIILEC 37

RESULT 15
US-10-367-057-16
Sequence 16, Application US/10367057
Publication No. US20050100554A1
GENERAL INFORMATION:
APPLICANT: CUTHILL, SCOTT;
APPLICANT: JACKSON, AMANDA;
APPLICANT: LEWIN, DAVID A.;
APPLICANT: OOI, CHEAN ENG
TITLE OF INVENTION: Complexes and Methods of Using Same
FILE REFERENCE: 21402-559
CURRENT APPLICATION NUMBER: US/10/367,057
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: 60/256,911
PRIOR FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 198
SOFTWARE: Curaseqlet version 0.1
SEQ ID NO 16
LENGTH: 158

TYPE: PRT
ORGANISM: Homo sapiens
US-10-367-057-16

Query Match 100.0%; Score 49; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD1ILEC 9
Db 29 T1HD1ILEC 37

RESULT 16
US-11-021-949-13

Sequence 13, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SAMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
TITLE OF INVENTION: AND METHODS OF THEIR USE
FILE REFERENCE: VITA-012
CURRENT FILING DATE: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 13
LENGTH: 158
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
US-11-021-949-13

Query Match 100.0%; Score 49; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD1ILEC 9
Db 29 T1HD1ILEC 37

RESULT 17

US-10-472-724-2

Sequence 2, Application US/10472724
Publication No. US20040171806A1
GENERAL INFORMATION:
APPLICANT: Cid-Arregui, Angel
APPLICANT: Zur Hausen, Harald
TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
FILE REFERENCE: 4121-154
CURRENT APPLICATION NUMBER: US/10/472,724
CURRENT FILING DATE: 2003-09-17
PRIOR APPLICATION NUMBER: PCT/EP02/03271
PRIOR FILING DATE: 2002-03-22
PRIOR APPLICATION NUMBER: EP 01107271.7
PRIOR FILING DATE: 2001-03-23
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn version 3.2
SEQ ID NO 2
LENGTH: 171
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Synthetic Construct
US-10-472-724-2

Query Match 100.0%; Score 49; DB 4; Length 171;

Best Local Similarity 100.0%; Pred. No. 0.23;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD1ILEC 9
Db 34 T1HD1ILEC 42

RESULT 18

US-11-072-288-1

Sequence 1, Application US/11072288
Publication No. US20050159386A1
GENERAL INFORMATION:
APPLICANT: KIENY, Marie-Paule
APPLICANT: BALLOU, Jean-Marc
APPLICANT: BIZOUANE, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
FILE REFERENCE: 017753-122
CURRENT APPLICATION NUMBER: US/11/072,288
CURRENT FILING DATE: 2005-03-07
PRIOR APPLICATION NUMBER: US/09/462,993
PRIOR FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: PCT/FR98/01576
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: PatentIn Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-11-072-288-1

Query Match 100.0%; Score 49; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.34;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD1ILEC 9
Db 57 T1HD1ILEC 65

RESULT 19

US-09-367-309A-1

Sequence 1, Application US/09367309A
Publication No. US20020081329A1
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JTM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 49; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.37;

Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDI1LEC 9
Db 29 T1HDI1LEC 37

RESULT 20

US-10-000-903-4
; Sequence 4, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernandez
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-4

Query Match 100.0%; Score 49; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.38; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 T1HDI1LEC 9
Db 135 T1HDI1LEC 143

RESULT 21

US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: Influenzae B and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4

Query Match 100.0%; Score 49; DB 5; Length 273;

Best Local Similarity 100.0%; Pred. No. 0.38; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 T1HDI1LEC 9
Db 135 T1HDI1LEC 143

RESULT 22

US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernandez
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-10

Query Match 100.0%; Score 49; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.41; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 T1HDI1LEC 9
Db 154 T1HDI1LEC 162

RESULT 23

US-10-899-771-10
; Sequence 10, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-10

Query Match 100.0%; Score 49; DB 5; Length 292;

Query Match 100.0%; Score 49; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.41;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 154 T1HDIILEC 162

RESULT 24

US-10-000-903-6
; Sequence 6, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000.903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-6

Query Match 100.0%; Score 49; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.53;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 135 T1HDIILEC 143

RESULT 25

US-10-899-771-6
; Sequence 6, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899.771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581.976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeraic protein (protein D from Haemophilus
; OTHER INFORMATION: Influenzae B and B67 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-6

Query Match 100.0%; Score 49; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.53;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 135 T1HDIILEC 143

RESULT 26

US-10-000-903-14
; Sequence 14, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000.903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-14

Query Match 100.0%; Score 49; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HDIILEC 9
Db 154 T1HDIILEC 162

RESULT 27

US-10-899-771-14
; Sequence 14, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899.771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581.976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeraic protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-14

US-10-899-771-14

Query Match 100.0%; Score 49; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDILLEC 9
|||||
Db 154 TTHDILLEC 162

RESULT 28

US-10-367-095-10
; Sequence 10, Application US/10367095
; Publication No. US20030228696A1

GENERAL INFORMATION:

APPLICANT: Robin A. Robinson

TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line

FILE REFERENCE: 44149-1US1

CURRENT APPLICATION NUMBER: US/10/367,095

PRIOR FILING DATE: 2003-02-14

PRIOR APPLICATION NUMBER: US 60/356,119

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,161

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,118

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,133

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,157

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,156

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,123

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,135

PRIOR FILING DATE: 2002-02-14

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FaastSeq for Windows Version 4.0

SEQ ID NO 10

LENGTH: 536

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: HPV-16 L2/E6 fusion protein

US-10-367-095-10

Query Match 100.0%; Score 49; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 0.79;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDILLEC 9
|||||
Db 499 TTHDILLEC 507

RESULT 29

US-10-368-046-10

; Sequence 10, Application US/10368046

; Publication No. US20040063188A1

GENERAL INFORMATION:

APPLICANT: Robin A. Robinson

TITLE OF INVENTION: Method for Isolation and Purification of

FILE REFERENCE: 44149-3US1

CURRENT APPLICATION NUMBER: US/10/368,046

CURRENT FILING DATE: 2003-02-15

PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,161

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,118

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,133

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,157

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,156

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,123

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,113

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,154

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,135

PRIOR FILING DATE: 2002-02-14

Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FaastSeq for Windows Version 4.0

SEQ ID NO 10

LENGTH: 536

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: HPV-16 L2/E6 fusion protein

US-10-368-046-10

QY 1 TTHDILLEC 9
|||||
Db 499 TTHDILLEC 507

RESULT 30

US-10-367-367-10

; Sequence 10, Application US/10367367

; Publication No. US20040121465A1

GENERAL INFORMATION:

APPLICANT: Robin A. Robinson

TITLE OF INVENTION: Optimization of Gene Sequences of

FILE REFERENCE: 44149-2US1

CURRENT APPLICATION NUMBER: US/10/367,367

CURRENT FILING DATE: 2003-02-15

PRIOR APPLICATION NUMBER: US 60/356,119

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,161

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,118

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,133

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,157

PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,156

PRIOR FILING DATE: 2002-02-14

NUMBER OF SEQ ID NOS: 13

SOFTWARE: FaastSeq for Windows Version 4.0

SEQ ID NO 10

LENGTH: 536

TYPE: PRT

ORGANISM: Artificial Sequence

FEATURE:

OTHER INFORMATION: HPV-16 L2/E6 fusion protein

US-10-367-367-10

Query Match 100.0%; Score 49; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 0.79; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0;

Qy 1 THDIILEC 9
Db 499 THDIILEC 507

RESULT 31

US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:
; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 19065/2132
; CURRENT APPLICATION NUMBER: US/10/918,337
; PRIOR FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-918-337-10

Query Match 100.0%; Score 49; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 0.79; Mismatches 0; Indels 0; Gaps 0;
Matches 9; Conservative 0;

Qy 1 THDIILEC 9
Db 499 THDIILEC 507

RESULT 32

US-10-476-570-10
; Sequence 10, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERS, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof

FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 10
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 30-50
US-10-476-570-10

Query Match 89.8%; Score 44; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.22; Mismatches 0; Indels 0; Gaps 0;
Matches 8; Conservative 0;

Qy 2 IHDIILEC 9
Db 1 IHDIILEC 8

RESULT 33

US-10-437-963-149433
; Sequence 149433, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yinhua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 149433
; LENGTH: 306
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_49767C.1.pep
US-10-437-963-149433

Query Match 83.7%; Score 41; DB 4; Length 306;
Best Local Similarity 66.7%; Pred. No. 15; Mismatches 1; Indels 0; Gaps 0;
Matches 6; Conservative 2;

Qy 1 THDIILEC 9
Db 192 TVHDIAIEC 200

RESULT 34

US-10-476-570-24
; Sequence 24, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERS, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard

;; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
;; FILE REFERENCE: 45636-5071-US
;; CURRENT APPLICATION NUMBER: US/10/476,570
;; PRIOR FILING DATE: 2003-11-04
;; PRIOR APPLICATION NUMBER: PCT/FR02/01533
;; PRIOR FILING DATE: 2002-05-03
;; PRIOR APPLICATION NUMBER: FR 01 05980
;; NUMBER OF SEQ ID NOS: 63
;; SOFTWARE: PatentIn Ver. 2.1
;; SEQ ID NO 24
;; LENGTH: 15
;; TYPE: PRT
;; ORGANISM: artificial sequence
;; FEATURE:
;; OTHER INFORMATION: Description of the artificial sequence: peptide E6 31-45
US-10-476-570-24

Query Match 81.6%; Score 40; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.89;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIILEC 9
Db 1 HDIILEC 7

RESULT 35
US-10-437-963-153797
;; Sequence 153797, Application US/10437963
;; Publication No. US2004012343A1
;; GENERAL INFORMATION:
;; APPLICANT: La Rosa, Thomas J.
;; APPLICANT: Kovalic, David K.
;; APPLICANT: Zhou, Yihua
;; APPLICANT: Cao, Yongwei
;; APPLICANT: Wu, Wei
;; APPLICANT: Boukharov, Andrey A.
;; APPLICANT: Barbazuk, Brad
;; APPLICANT: Li, Ping
;; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated with
;; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
;; FILE REFERENCE: 38-21(53221)B
;; CURRENT APPLICATION NUMBER: US/10/437,963
;; CURRENT FILING DATE: 2003-05-14
;; NUMBER OF SEQ ID NOS: 204966
;; SEQ ID NO 153797
;; LENGTH: 150
;; TYPE: PRT
;; ORGANISM: Oryza sativa
;; NAME/KEY: unsure
;; LOCATION: (1)..(150)
;; OTHER INFORMATION: unsure at all Xaa locations
;; FEATURE:
;; OTHER INFORMATION: Clone ID: PAT_MRT4530_53718C.1.pcp
US-10-437-963-153797

Query Match 81.6%; Score 40; DB 4; Length 150;
Best Local Similarity 66.7%; Pred. No. 11;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 THDIILEC 9
Db 121 TVHDIIVEC 129

RESULT 36
US-10-424-599-275411
;; Sequence 275411, Application US/10424599
;; Publication No. US20040031072A1
;; GENERAL INFORMATION:

;; APPLICANT: La Rosa, Thomas J.
;; APPLICANT: Kovalic, David K.
;; APPLICANT: Zhou, Yihua
;; APPLICANT: Cao, Yongwei
;; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated with
;; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
;; FILE REFERENCE: 38-21(53223)B
;; CURRENT APPLICATION NUMBER: US/10/424,599
;; CURRENT FILING DATE: 2003-04-28
;; NUMBER OF SEQ ID NOS: 285684
;; SEQ ID NO 275411
;; LENGTH: 297
;; TYPE: PRT
;; ORGANISM: Glycine max
;; FEATURE:
;; OTHER INFORMATION: Clone ID: PAT_MRT3847_90717C.1.pcp
US-10-424-599-275411

Query Match 79.6%; Score 39; DB 4; Length 297;
Best Local Similarity 62.5%; Pred. No. 35;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IHDIILEC 9
Db 179 IHDLVIEC 186

RESULT 37
US-10-424-599-168507
;; Sequence 168507, Application US/10424599
;; Publication No. US20040031072A1
;; GENERAL INFORMATION:
;; APPLICANT: La Rosa, Thomas J.
;; APPLICANT: Kovalic, David K.
;; APPLICANT: Zhou, Yihua
;; APPLICANT: Cao, Yongwei
;; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated with
;; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
;; FILE REFERENCE: 38-21(53223)B
;; CURRENT APPLICATION NUMBER: US/10/424,599
;; CURRENT FILING DATE: 2003-04-28
;; NUMBER OF SEQ ID NOS: 285684
;; SEQ ID NO 168507
;; LENGTH: 298
;; TYPE: PRT
;; ORGANISM: Glycine max
;; NAME/KEY: unsure
;; LOCATION: (1)..(298)
;; OTHER INFORMATION: unsure at all Xaa locations
;; FEATURE:
;; OTHER INFORMATION: Clone ID: PAT_MRT3847_123177C.1.pcp
US-10-424-599-168507

Query Match 79.6%; Score 39; DB 4; Length 298;
Best Local Similarity 62.5%; Pred. No. 35;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 IHDIILEC 9
Db 178 IHDLVIEC 185

RESULT 38
US-10-732-923-21847
;; Sequence 21847, Application US/10732923
;; Publication No. US20050108791A1
;; GENERAL INFORMATION:
;; APPLICANT: Edgerton, Michael D
;; TITLE OF INVENTION: TRANSGENIC PLANTS WITH IMPROVED PHENOTYPES
;; FILE REFERENCE: 38-15(52796)C
;; CURRENT APPLICATION NUMBER: US/10/732,923
;; CURRENT FILING DATE: 2003-12-10

```
; PRIOR APPLICATION NUMBER: 10/310,154
; PRIOR FILING DATE: 2002-12-04
; NUMBER OF SEQ ID NOS: 24149
; SEQ ID NO 21847
; LENGTH: 145
; TYPE: PRT
; ORGANISM: Bacillus halodurans
US-10-732-923-21847

Query Match      75.5%; Score 37; DB 5; Length 145;
Best Local Similarity 66.7%; Pred. No. 40;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 THDIILEC 9
Db      46 TDHVTILEC 54

RESULT 39
US-11-021-949-361
; Sequence 361, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FASTSEQ for Windows Version 4.0
; SEQ ID NO 361
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-361

Query Match      75.5%; Score 37; DB 6; Length 158;
Best Local Similarity 55.6%; Pred. No. 43;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 THDIILEC 9
Db      24 TLHDITDC 32

RESULT 40
US-10-739-930-10846
; Sequence 10846, Application US/10739930
; Publication No. US20040216190A1
; GENERAL INFORMATION:
; APPLICANT: KOVALIC, DAVID K.
; TITLE OF INVENTION: NOCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
; FILE REFERENCE: 38-21(53377)B
; CURRENT APPLICATION NUMBER: US/10/739,930
; CURRENT FILING DATE: 2003-12-18
; NUMBER OF SEQ ID NOS: 11088
; SEQ ID NO 10846
; LENGTH: 554
; TYPE: PRT
; ORGANISM: Triticum aestivum
; NAME/KEY: unsure
; LOCATION: (1)..(554)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:

; OTHER INFORMATION: Clone ID: TRIAE-23APR03-C7327_1.P
US-10-739-930-10846

Query Match      75.5%; Score 37; DB 5; Length 554;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      2 IHDIILEC 9
Db      458 VHDVTILEC 465

RESULT 41
US-10-029-386-33593
; Sequence 33593, Application US/10029386
; Publication No. US20030194704A1
; GENERAL INFORMATION:
; APPLICANT: Penn, Sharon G.
; APPLICANT: Rank, David R.
; APPLICANT: Hanzel, David K.
; TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR GI
; TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
; FILE REFERENCE: AROMICA-X-2
; CURRENT APPLICATION NUMBER: US/10/029,386
; CURRENT FILING DATE: 2001-12-20
; NUMBER OF SEQ ID NOS: 34288
; SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
; SEQ ID NO 33593
; LENGTH: 70
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; OTHER INFORMATION: MAP TO CHR9.1
; OTHER INFORMATION: EXPRESSED IN PETAL LIVER, SIGNAL = 1.1
; OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.59
; OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.79
; OTHER INFORMATION: SWISSPROT HIT: Q92574, EVALUE 2.00e-30
US-10-029-386-33593

Query Match      73.5%; Score 36; DB 4; Length 70;
Best Local Similarity 57.1%; Pred. No. 28;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY      3 HDIILEC 9
Db      16 HDVTILEC 22

RESULT 42
US-10-425-115-205547
; Sequence 205547, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: KOVALIC, DAVID K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53322)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 205547
; LENGTH: 83
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(83)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_119045C.1.pep
```

US-10-425-115-205547

Query Match 73.5%; Score 36; DB 4; Length 83;
Best Local Similarity 75.0%; Pred. No. 34;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 THDILLEC 9
|:|:|:|:
Db 53 IHGVILLEC 60

RESULT 43

US-10-425-115-194068
; Sequence 194068, Application US/10425115
; Publication NO. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 194068
; LENGTH: 117
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)-(117)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_108570C.1.pep
US-10-425-115-194068

Query Match 73.5%; Score 36; DB 4; Length 117;
Best Local Similarity 55.6%; Pred. No. 49;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 THDILLEC 9
|:|:|:|:
Db 21 TLHDIMMAC 29

RESULT 44

US-11-021-949-31
; Sequence 31, Application US/11021949
; Publication NO. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 31
; LENGTH: 162
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-31

Query Match 73.5%; Score 36; DB 6; Length 162;

Best Local Similarity 44.4%; Pred. No. 70;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

QY 1 THDILLEC 9
|:|:|:|:
Db 28 TLHDVITDC 36

RESULT 45

US-10-425-115-358720
; Sequence 358720, Application US/10425115
; Publication NO. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 358720
; LENGTH: 164
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_90319C.1.pep
US-10-425-115-358720

Query Match 73.5%; Score 36; DB 4; Length 164;
Best Local Similarity 75.0%; Pred. No. 70;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 THDILLEC 8
|:|:|:|:
Db 82 TLHDILB 89

RESULT 46

US-10-425-115-358724
; Sequence 358724, Application US/10425115
; Publication NO. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 358724
; LENGTH: 173
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_90322C.1.pep
US-10-425-115-358724

Query Match 73.5%; Score 36; DB 4; Length 173;
Best Local Similarity 75.0%; Pred. No. 75;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 THDILLEC 8
|:|:|:|:
Db 82 TLHDILB 89

RESULT 47

US-10-947-979-18
; Sequence 18, Application US/10947979
; Publication No. US20050102717A1
; GENERAL INFORMATION:
; APPLICANT: Altier, Daniel
; APPLICANT: Bao, Zhongmeng
; APPLICANT: Lu, Guihua
; APPLICANT: Navarro-Acevedo, Pedro
; APPLICANT: Sewalt, Vincent
; APPLICANT: Simmons, Carl
; APPLICANT: Yalpani, Nasser
; TITLE OF INVENTION: Crop Plant Cystatin Proteinase
; TITLE OF INVENTION: Inhibitors and Their Use
; FILE REFERENCE: 1498
; CURRENT APPLICATION NUMBER: US/10/947,979
; CURRENT FILING DATE: 2004-09-23
; PRIOR APPLICATION NUMBER: 60/505,948
; PRIOR FILING DATE: 2004-09-25
; NUMBER OF SEQ ID NOS: 195
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 18
; LENGTH: 174
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (50)...(59)
; OTHER INFORMATION: N-terminal alpha-1 helix domain
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (77)...(87)
; OTHER INFORMATION: First hairpin loop domain
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (100)...(108)
; OTHER INFORMATION: Second hairpin loop domain
US-10-947-979-18

Query Match 73.5%; Score 36; DB 5; Length 174;
Best Local Similarity 75.0%; Pred. No. 76;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TINDLLE 8
|:|:|:|:|
Db 82 TINDLLE 89

RESULT 48
US-10-425-115-236213
; Sequence 236213, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 236213
; LENGTH: 176
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: MWT4577_147007C.1.pep
US-10-425-115-236213

Query Match 73.5%; Score 36; DB 4; Length 176;
Best Local Similarity 75.0%; Pred. No. 76;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TINDLLE 8
|:|:|:|:|
Db 85 TINDLLE 92

RESULT 49
US-10-947-979-10
; Sequence 10, Application US/10947979
; Publication No. US20050102717A1
; GENERAL INFORMATION:
; APPLICANT: Altier, Daniel
; APPLICANT: Bao, Zhongmeng
; APPLICANT: Lu, Guihua
; APPLICANT: Navarro-Acevedo, Pedro
; APPLICANT: Sewalt, Vincent
; APPLICANT: Simmons, Carl
; APPLICANT: Yalpani, Nasser
; TITLE OF INVENTION: Crop Plant Cystatin Proteinase
; TITLE OF INVENTION: Inhibitors and Their Use
; FILE REFERENCE: 1498
; CURRENT APPLICATION NUMBER: US/10/947,979
; CURRENT FILING DATE: 2004-09-23
; PRIOR APPLICATION NUMBER: 60/505,948
; PRIOR FILING DATE: 2004-09-25
; NUMBER OF SEQ ID NOS: 195
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 176
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (53)...(62)
; OTHER INFORMATION: N-terminal alpha-1 helix domain
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (80)...(90)
; OTHER INFORMATION: First hairpin loop domain
; FEATURE:
; NAME/KEY: DOMAIN
; LOCATION: (103)...(111)
; OTHER INFORMATION: Second hairpin loop domain
US-10-947-979-10

Query Match 73.5%; Score 36; DB 5; Length 176;
Best Local Similarity 75.0%; Pred. No. 76;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Oy 1 TINDLLE 8
|:|:|:|:|
Db 85 TINDLLE 92

RESULT 50
US-10-425-114-56403
; Sequence 56403, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53113)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 56403
; LENGTH: 211

```

; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: UC-ZMFLB73208D10_FLI.pep
US-10-425-114-56403

```

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Query Match      73.5%; Score 36; DB 4; Length 211;
Best Local Similarity 75.0%; Pred. No. 93;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

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```

OY      1 THDIIIL 8
        |:|:|:|
Db      120 THDIIIL 127

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Job time : 60.8 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:08:06 ; Search time 8.4 Seconds
(without alignments)
49,591 Million cell updates/sec

Title: US-08-170-344-7
Perfect score: 49
Sequence: 1 T1HD1LTC 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries

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SUMMARIES

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2	49	100.0	158	11 US-11-206-138-3	Sequence 3, Appl1
3	49	100.0	248	9 US-10-530-253-1	Sequence 1, Appl1
4	49	100.0	248	9 US-10-530-253-3	Sequence 3, Appl1
5	49	100.0	248	9 US-10-530-253-5	Sequence 5, Appl1
6	49	100.0	248	9 US-10-530-253-9	Sequence 7, Appl1
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10	36	73.5	158	9 US-10-530-253-26	Sequence 26, Appl1
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12	36	73.5	298	11 US-11-172-740-1141	Sequence 1141, Ap
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16	35	71.4	220	11 US-11-087-099-4165	Sequence 4165, Ap
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253	29	59.2	554	11	US-11-098-686-10351	Sequence 10351, A	326	28	57.1	336	9	US-10-873-528-1850	Sequence 8640, App
254	29	59.2	586	9	US-10-500-941-4	Sequence 4, App1	327	28	57.1	336	11	US-11-188-298-8640	Sequence 12800, A
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262	29	59.2	739	9	US-10-505-263-96	Sequence 96, App1	335	28	57.1	378	11	US-11-096-568A-8228	Sequence 8228, App
263	29	59.2	756	11	US-11-188-298-6623	Sequence 6623, App	336	28	57.1	380	9	US-10-330-773-443	Sequence 443, App
264	29	59.2	780	9	US-10-878-556A-197	Sequence 197, App	337	28	57.1	382	11	US-11-096-568A-15422	Sequence 15422, A
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267	29	59.2	780	9	US-10-784-004-462	Sequence 462, App	340	28	57.1	398	8	US-10-511-937-2405	Sequence 2405, App
268	29	59.2	780	9	US-10-784-004-779	Sequence 779, App	341	28	57.1	398	9	US-10-821-234-1583	Sequence 1583, App
269	29	59.2	780	9	US-10-784-004-780	Sequence 780, App	342	28	57.1	405	11	US-11-096-568A-7201	Sequence 7201, App
270	29	59.2	838	11	US-11-188-298-3797	Sequence 3797, App	343	28	57.1	406	11	US-11-096-568A-7200	Sequence 7199, App
271	29	59.2	840	9	US-10-645-441-1	Sequence 1, App1	344	28	57.1	407	11	US-11-096-568A-7199	Sequence 7, App1
272	29	59.2	840	9	US-10-725-475-16	Sequence 16, App1	345	28	57.1	412	11	US-11-093-808-1	Sequence 8227, App
273	29	59.2	842	9	US-10-645-441-2	Sequence 2, App1	346	28	57.1	413	11	US-11-096-568A-15421	Sequence 15421, A
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285	29	59.2	1307	11	US-11-096-568A-30285	Sequence 7, App1	357	28	57.1	483	11	US-11-188-298-4164	Sequence 491, App
286	29	59.2	1382	11	US-11-208-414-7	Sequence 9, App1	358	28	57.1	496	11	US-11-188-298-4409	Sequence 28, App1
287	29	59.2	1382	11	US-11-208-414-9	Sequence 22, App1	359	28	57.1	499	9	US-10-330-773-491	Sequence 22316, A
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290	29	59.2	15	9	US-10-530-061-1735	Sequence 1735, App	362	28	57.1	503	11	US-11-188-298-22316	Sequence 230, App
291	29	59.2	88	11	US-11-098-686-11213	Sequence 11213, A	363	28	57.1	506	11	US-11-098-686-11213	Sequence 233, App
292	29	59.2	88	11	US-11-098-686-11213	Sequence 14086, A	364	28	57.1	508	11	US-11-098-686-11213	Sequence 24367, A
293	29	59.2	103	9	US-11-096-568A-14086	Sequence 14085, A	365	28	57.1	517	11	US-11-096-568A-34367	Sequence 14, App1
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295	29	59.2	103	9	US-10-506-454-452	Sequence 14292, A	367	28	57.1	527	11	US-11-096-568A-34367	Sequence 3870, App
296	29	59.2	103	9	US-10-506-454-452	Sequence 14292, A	368	28	57.1	542	11	US-11-096-568A-34367	Sequence 2956, App
297	29	59.2	103	9	US-10-506-454-452	Sequence 14292, A	369	28	57.1	550	11	US-11-096-568A-34367	Sequence 2956, App
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394	28	57.1	833	11	US-11-055-822-212	Sequence 212, App	467	27	55.1	227	9	US-10-467-657-2624	Sequence 2624, Ap
395	28	57.1	833	11	US-11-055-822-710	Sequence 710, App	468	27	55.1	229	11	US-11-100-183-46	Sequence 46, Appl
396	28	57.1	833	11	US-11-239-674-90	Sequence 90, Appl	469	27	55.1	233	11	US-11-096-568A-330962	Sequence 30962, A
397	28	57.1	897	11	US-11-072-512-2474	Sequence 2474, Ap	470	27	55.1	241	11	US-11-188-298-22369	Sequence 22369, A
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399	28	57.1	915	11	US-11-144-987-22	Sequence 22, Appl	472	27	55.1	256	11	US-11-188-298-3324	Sequence 3324, App
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403	28	57.1	917	11	US-11-144-987-20	Sequence 20, Appl	476	27	55.1	271	11	US-11-188-298-4998	Sequence 4948, Ap
404	28	57.1	917	11	US-11-144-987-24	Sequence 24, Appl	477	27	55.1	276	11	US-11-087-099-9241	Sequence 2221, Ap
405	28	57.1	917	11	US-11-144-987-26	Sequence 26, Appl	478	27	55.1	283	11	US-11-148-410-4	Sequence 4, Appl1
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418	28	57.1	998	11	US-11-203-251A-83	Sequence 83, Appl	491	27	55.1	335	11	US-11-182-946-7	Sequence 4573, Ap
419	28	57.1	998	11	US-11-203-251A-88	Sequence 88, Appl	492	27	55.1	335	11	US-11-096-568A-30960	Sequence 30960, A
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422	28	57.1	1044	11	US-11-091-668-2	Sequence 2, Appl1	495	27	55.1	335	11	US-11-188-298-8085	Sequence 8085, Ap
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425	28	57.1	1055	11	US-11-203-251A-86	Sequence 86, Appl	498	27	55.1	335	11	US-11-188-298-13038	Sequence 13038, A
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442	28	57.1	2333	11	US-11-096-281-13	Sequence 13, Appl	515	27	55.1	341	11	US-11-188-298-7585	Sequence 7585, Ap
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455	27.5	56.1	1366	9	US-11-043-693-34	Sequence 34, Appl	528	27	55.1	350	11	US-11-087-099-2497	Sequence 2497, Ap
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459	27	55.1	92	11	US-11-094-142-34	Sequence 6454, Ap	532	27	55.1	351	11	US-11-087-099-6490	Sequence 6490, Ap

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534	27	55.1	353	11	US-11-216-267-32	Sequence 32, Appl	607	27	55.1	825	11	US-11-188-298-13018	Sequence 13018, A
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536	27	55.1	361	11	US-11-096-568A-25811	Sequence 25811, A	609	27	55.1	855	11	US-11-188-298-20385	Sequence 20385, A
537	27	55.1	367	11	US-11-188-298-8631	Sequence 8631, Ap	610	27	55.1	920	11	US-11-188-298-12858	Sequence 12858, A
538	27	55.1	375	11	US-11-174-341-173	Sequence 173, App	611	27	55.1	979	11	US-10-330-773-376	Sequence 376, App
539	27	55.1	381	11	US-11-087-099-1428	Sequence 3428, App	612	27	55.1	1035	11	US-11-188-298-8170	Sequence 8170, Ap
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542	27	55.1	398	11	US-11-096-568A-1134	Sequence 1134, Ap	615	27	55.1	1036	9	US-10-243-116-104	Sequence 104, App
543	27	55.1	403	11	US-11-188-298-8075	Sequence 8075, Ap	616	27	55.1	1036	9	US-10-243-136-104	Sequence 104, App
544	27	55.1	404	11	US-11-096-568A-5191	Sequence 5191, Ap	617	27	55.1	1036	9	US-10-243-189-104	Sequence 104, App
545	27	55.1	412	11	US-11-093-808-9	Sequence 9, Appl1	618	27	55.1	1036	9	US-10-243-215-104	Sequence 104, App
546	27	55.1	412	11	US-11-093-808-10	Sequence 10, Appl1	619	27	55.1	1036	9	US-10-243-226-104	Sequence 104, App
547	27	55.1	412	11	US-11-093-808-11	Sequence 11, Appl1	620	27	55.1	1036	9	US-10-243-228-104	Sequence 104, App
548	27	55.1	412	11	US-11-093-808-12	Sequence 12, Appl1	621	27	55.1	1036	9	US-10-243-304-104	Sequence 104, App
549	27	55.1	412	11	US-11-093-808-13	Sequence 13, Appl1	622	27	55.1	1036	9	US-10-243-318-104	Sequence 104, App
550	27	55.1	412	11	US-11-093-808-14	Sequence 14, Appl1	623	27	55.1	1036	9	US-10-243-345-104	Sequence 104, App
551	27	55.1	412	11	US-11-087-099-499	Sequence 499, App	624	27	55.1	1036	9	US-10-243-357-104	Sequence 104, App
552	27	55.1	414	11	US-11-174-341-160	Sequence 160, App	625	27	55.1	1036	9	US-10-245-083-104	Sequence 104, App
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571	27	55.1	514	11	US-11-188-298-21625	Sequence 21625, A	644	27	55.1	2305	11	US-11-126-313-33	Sequence 1002, Ap
572	27	55.1	515	9	US-10-915-002-240	Sequence 240, App	645	27	55.1	4834	8	US-10-453-372-1002	Sequence 827, App
573	27	55.1	523	11	US-11-087-099-1873	Sequence 6809, Ap	646	27	55.1	4834	11	US-10-505-928-827	Sequence 20, Appl
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587	27	55.1	567	11	US-11-072-512-197	Sequence 197, App	660	26	53.1	125	7	US-09-978-360A-743	Sequence 224, App
588	27	55.1	593	9	US-10-330-773-116	Sequence 116, App	661	26	53.1	125	9	US-10-475-075-224	Sequence 499, App
589	27	55.1	616	11	US-11-188-298-17292	Sequence 17292, A	662	26	53.1	125	9	US-10-475-075-499	Sequence 967, App
590	27	55.1	623	9	US-10-793-626-1068	Sequence 1068, Ap	663	26	53.1	133	9	US-10-821-234-967	Sequence 8289, Ap
591	27	55.1	624	9	US-10-506-454-544	Sequence 544, App	664	26	53.1	134	11	US-11-096-568A-20906	Sequence 20906, A
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593	27	55.1	641	9	US-10-821-234-1519	Sequence 1519, App	666	26	53.1	135	9	US-11-087-099-8289	Sequence 3170, Ap
594	27	55.1	641	9	US-10-491-096-189	Sequence 189, App	667	26	53.1	153	11	US-11-096-568A-20905	Sequence 20905, A
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596	27	55.1	666	9	US-10-330-773-118	Sequence 118, App	669	26	53.1	158	11	US-11-087-099-3151	Sequence 2643, Ap
597	27	55.1	661	11	US-11-072-512-2369	Sequence 2369, App	670	26	53.1	159	11	US-11-096-568A-2643	Sequence 11569, Ap
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599	27	55.1	669	11	US-11-072-512-2373	Sequence 2373, App	672	26	53.1	175	11	US-11-096-568A-11569	Sequence 1588, Ap
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681	26	53.1	191	11	US-11-172-740-1734	Sequence 1734, Ap	754	26	53.1	323	11	US-11-098-686-10772	Sequence 10272, A
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684	26	53.1	192	11	US-11-172-740-1740	Sequence 1740, Ap	757	26	53.1	327	9	US-10-506-454-668	Sequence 866, App
685	26	53.1	192	11	US-11-172-740-1741	Sequence 1741, Ap	758	26	53.1	327	9	US-10-134-487-422	Sequence 422, App
686	26	53.1	192	11	US-11-172-740-1742	Sequence 1742, Ap	759	26	53.1	327	9	US-10-135-883-422	Sequence 422, App
687	26	53.1	192	11	US-11-172-740-1743	Sequence 1743, Ap	760	26	53.1	327	9	US-10-135-888-422	Sequence 422, App
688	26	53.1	192	11	US-11-172-740-1744	Sequence 1744, Ap	761	26	53.1	327	9	US-10-135-889-422	Sequence 422, App
689	26	53.1	196	11	US-11-172-740-2213	Sequence 2213, Ap	762	26	53.1	327	9	US-10-218-784-142	Sequence 142, App
690	26	53.1	196	11	US-11-172-740-2214	Sequence 2214, Ap	763	26	53.1	327	9	US-10-219-061-142	Sequence 142, App
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692	26	53.1	202	11	US-11-045-004-2660	Sequence 2660, Ap	765	26	53.1	327	9	US-10-219-064-142	Sequence 142, App
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707	26	53.1	233	11	US-11-096-568A-14614	Sequence 14614, A	780	26	53.1	335	11	US-11-188-298-2165	Sequence 21765, A
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714	26	53.1	250	11	US-11-096-568A-10833	Sequence 10833, A	787	26	53.1	341	9	US-10-793-626-3248	Sequence 3248, Ap
715	26	53.1	251	11	US-11-096-568A-22675	Sequence 22675, A	788	26	53.1	341	9	US-10-793-626-3248	Sequence 3248, Ap
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731	26	53.1	284	11	US-11-079-463-6467	Sequence 6467, Ap	804	26	53.1	348	11	US-11-188-298-17868	Sequence 17868, A
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736	26	53.1	295	11	US-11-188-298-12185	Sequence 12185, A	809	26	53.1	353	11	US-11-116-881A-405	Sequence 490, App
737	26	53.1	297	11	US-11-096-568A-10831	Sequence 10831, A	810	26	53.1	354	9	US-10-467-657-60	Sequence 60, App1
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742	26	53.1	302	11	US-11-188-298-13104	Sequence 13104, A	815	26	53.1	357	11	US-11-087-099-7487	Sequence 7487, Ap
743	26	53.1	308	11	US-11-087-099-7198	Sequence 7198, Ap	816	26	53.1	357	11	US-11-096-568A-28712	Sequence 28712, A
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746	26	53.1	313	11	US-11-188-298-274	Sequence 274, App	819	26	53.1	363	11	US-11-087-099-9262	Sequence 9262, Ap
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982 26 53.1 3623 9 US-10-955-561-593 Sequence 593, App
983 25.5 52.0 170 11 US-11-087-039-9220 Sequence 2220, App
984 25.5 52.0 212 11 US-11-098-686-11253 Sequence 11253, App
985 25.5 52.0 355 11 US-11-098-686-11427 Sequence 11427, App
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988 25.5 52.0 730 9 US-10-821-234-1019 Sequence 1019, App
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991 25.5 52.0 1062 11 US-11-124-367A-387 Sequence 387, App
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996 25 51.0 29 9 US-10-485-517-375 Sequence 375, App
997 25 51.0 37 9 US-10-729-131-22 Sequence 22, App
998 25 51.0 39 11 US-11-285-537-22 Sequence 22, App
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ALIGNMENTS

RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13

Query Match 100.0%; Score 49; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.015;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1

; GENERAL INFORMATION:
; APPLICANT: Healthbanc Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; CURRENT FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3

Query Match 100.0%; Score 49; DB 11; Length 158;
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Qy 1 TTHDIIIEC 9
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RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
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; ORGANISM: Human papillomavirus type 16
US-10-530-253-1

Query Match 100.0%; Score 49; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TTHDIIIEC 9
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RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929

;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 3
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;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-3

Query Match 100.0%; Score 49; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
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Db 22 TTHDIIIEC 30

RESULT 5
US-10-530-253-5
;; Sequence 5, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Cassetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M137-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; PRIOR FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02
;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
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;; LENGTH: 248
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 49; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
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Db 22 TTHDIIIEC 30

RESULT 6
US-10-530-253-7
;; Sequence 7, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Cassetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; APPLICANT: Susan P. McElhinney
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M137-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; PRIOR FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02
;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
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;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match 100.0%; Score 49; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
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Db 119 TTHDIIIEC 127

RESULT 7
US-10-530-253-9
;; Sequence 9, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Cassetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; APPLICANT: Susan P. McElhinney
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M137-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; PRIOR FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02
;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 9
;; LENGTH: 248
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 49; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 TTHDIIIEC 9
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Db 119 TTHDIIIEC 127

RESULT 8
US-10-530-253-11
;; Sequence 11, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Cassetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; APPLICANT: Susan P. McElhinney
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M137-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; PRIOR FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02
;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 11
;; LENGTH: 248
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 49; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD11EC 9
Db 119 T1HD11EC 127

RESULT 9

US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XIAOWU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192,923A
; PRIOR FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 49; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.025;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 T1HD11EC 9
Db 127 T1HD11EC 135

RESULT 10

US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 26
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 68
US-10-530-253-26

Query Match 73.5%; Score 36; DB 9; Length 158;
Best Local Similarity 44.4%; Pred. No. 5.9;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Qy 1 T1HD11EC 9
Db 24 T1HDV1DC 32

RESULT 11
US-11-172-740-1143

; Sequence 1143, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
; APPLICANT: MASCIA, Peter
; APPLICANT: ALEXANDROV, Nikolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
; FILE REFERENCE: 2750-1602PUS2
; CURRENT APPLICATION NUMBER: US/11/172,740
; PRIOR FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,800
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1143
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Zea mays

FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(202)
OTHER INFORMATION: Public GI no. 34903270
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for increasing chlorophyll and photosynthetic cap
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making lethal plants for genetic confinement
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making ornamental plants with modified flower
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making ornamental plants with modified leaves
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making plants with altered leaf shape eg curl
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making plants with increased biomass
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Uclity: Useful for making smaller plants
US-11-172-740-1143

Query Match 73.5%; Score 36; DB 11; Length 202;
Best Local Similarity 62.5%; Pred. No. 7.6;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 2 T1HD11EC 9
Db 42 VHD1FLDC 49

RESULT 12

US-11-172-740-1141
; Sequence 1141, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:

```

; APPLICANT: MASCIA, Peter
; APPLICANT: ALEXANDROV, Nickolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
; TITLE OF INVENTION: PLANT CHARACTERISTICS AND PHENOTYPES
; FILE REFERENCE: 2750-1602PUS2
; CURRENT FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,800
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1141
; LENGTH: 298
; TYPE: PRT
; ORGANISM: Triticum aestivum
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(298)
; OTHER INFORMATION: Ceres CLONE ID no. 788296
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; US-11-172-740-1141
; OTHER INFORMATION: Utility: Useful for making smaller plants

Query Match      73.5%; Score 36; DB 11; Length 298;
Best Local Similarity 62.5%; Pred. No. 11;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      2 INDILEC 9
DB      138 VHDIFLDC 145

RESULT 13
US-11-172-740-1144
; Sequence 1144, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
; APPLICANT: MASCIA, Peter
; APPLICANT: ALEXANDROV, Nickolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
; TITLE OF INVENTION: PLANT CHARACTERISTICS AND PHENOTYPES
; FILE REFERENCE: 2750-1602PUS2

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; CURRENT APPLICATION NUMBER: US/11/172,740
; CURRENT FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,800
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1144
; LENGTH: 306
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(306)
; OTHER INFORMATION: Public GI no. 56783703
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; US-11-172-740-1144
; OTHER INFORMATION: Utility: Useful for making smaller plants

Query Match      73.5%; Score 36; DB 11; Length 306;
Best Local Similarity 62.5%; Pred. No. 12;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      2 INDILEC 9
DB      146 VHDIFLDC 153

RESULT 14
US-11-172-740-1142
; Sequence 1142, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
; APPLICANT: MASCIA, Peter
; APPLICANT: ALEXANDROV, Nickolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
; TITLE OF INVENTION: PLANT CHARACTERISTICS AND PHENOTYPES
; FILE REFERENCE: 2750-1602PUS2
; CURRENT FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30

```



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FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 6614
LENGTH: 301
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)-(301)
OTHER INFORMATION: Ceres Seq. ID no. 14315990
US-11-096-568A-6614

Query Match      71.4%; Score 35; DB 11; Length 301;
Best Local Similarity 50.0%; Pred. No. 18;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      2 IHDIIIEC 9
      :||: |||
Db      181 VHDVITIC 188

RESULT 19
US-11-031-206-106
Sequence 106, Application US/11031206
Publication No. US20060031959A1
GENERAL INFORMATION:
APPLICANT: Rafalski, Antoni
APPLICANT: Miao, Guo-Hua
APPLICANT: Falco, Saverio Carl
APPLICANT: Sakai, Hajime
APPLICANT: Pamodu, Omolayo O.
APPLICANT: Odell, Joan T.
APPLICANT: Meyers, Blake
APPLICANT: Thorpe, Catherine
APPLICANT: Weng, Zude
TITLE OF INVENTION: Nucleic Acid Fragments Encoding Proteins Involved in
TITLE OF INVENTION: Stress Response
FILE REFERENCE: BR157 US NA
CURRENT APPLICATION NUMBER: US/11/031,206
CURRENT FILING DATE: 2005-01-07
PRIOR APPLICATION NUMBER: US/09/566,394
PRIOR FILING DATE: 2000-05-05
PRIOR APPLICATION NUMBER: 60/133038
PRIOR FILING DATE: 1999-05-07
PRIOR APPLICATION NUMBER: 60/133042
PRIOR FILING DATE: 1999-05-07
PRIOR APPLICATION NUMBER: 60/133427
PRIOR FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: 60/133437
PRIOR FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: 60/133428
PRIOR FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: 60/133438
PRIOR FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: 60/133436
PRIOR FILING DATE: 1999-05-11
PRIOR APPLICATION NUMBER: 60/137667
PRIOR FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 208
SOFTWARE: Microsoft Office 97
SEQ ID NO 106
LENGTH: 527
TYPE: PRT
ORGANISM: Zea mays
US-11-031-206-106

Query Match      71.4%; Score 35; DB 11; Length 527;
Best Local Similarity 75.0%; Pred. No. 33;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

QY 1 TIIIDILE 8

Db 379 TIIIDILE 386

```
RESULT 20
US-10-530-061-53
Sequence 53, Application US/10530061
Publication No. US20060079453A1
GENERAL INFORMATION:
APPLICANT: SIDNEY, JOHN
APPLICANT: SOUTHWOOD, SCOTT
APPLICANT: SETTE, ALESSANDRO
TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
FILE REFERENCE: 2060.03US02/EKS/M-M
CURRENT APPLICATION NUMBER: US/10/530,061
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31308
PRIOR FILING DATE: 2003-10-03
PRIOR APPLICATION NUMBER: 60/416,207
PRIOR FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/417,269
PRIOR FILING DATE: 2002-10-08
NUMBER OF SEQ ID NOS: 2503
SOFTWARE: PatentIn version 3.3
SEQ ID NO 53
LENGTH: 10
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-530-061-53
```

Query Match 69.4%; Score 34; DB 9; Length 10;
Best Local Similarity 87.5%; Pred. No. 0.81;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 IHDIIIEC 9
 :|||: |||
Db 1 ITDIIIEC 8

```
RESULT 21
US-10-530-061-114
Sequence 114, Application US/10530061
Publication No. US20060079453A1
GENERAL INFORMATION:
APPLICANT: SIDNEY, JOHN
APPLICANT: SOUTHWOOD, SCOTT
APPLICANT: SETTE, ALESSANDRO
TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
FILE REFERENCE: 2060.03US02/EKS/M-M
CURRENT APPLICATION NUMBER: US/10/530,061
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US03/31308
PRIOR FILING DATE: 2003-10-03
PRIOR APPLICATION NUMBER: 60/416,207
PRIOR FILING DATE: 2002-10-03
PRIOR APPLICATION NUMBER: 60/417,269
PRIOR FILING DATE: 2002-10-08
NUMBER OF SEQ ID NOS: 2503
SOFTWARE: PatentIn version 3.3
SEQ ID NO 114
LENGTH: 10
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-530-061-114
```

Query Match 69.4%; Score 34; DB 9; Length 10;
Best Local Similarity 87.5%; Pred. No. 0.81;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 IHDIIIEC 9
 :|||: |||
Db 1 ITDIIIEC 8

RESULT 22
US-10-530-061-1673
; Sequence 1673, Application US/10530061
; Publication No. US20060079453a1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.03US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 1673
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-1673

Query Match 69.4%; Score 34; DB 9; Length 15;
Best Local Similarity 66.7%; Pred. No. 1.2;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 THHIIIEC 9
|||:|:|
Db 3 THHIIIEC 11

RESULT 23
US-10-530-253-17
; Sequence 17, Application US/10530253
; Publication No. US20060014926a1
; GENERAL INFORMATION:
; APPLICANT: Casagetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 17
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 33
US-10-530-253-17

Query Match 69.4%; Score 34; DB 9; Length 149;
Best Local Similarity 66.7%; Pred. No. 1.4;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 THHIIIEC 9
|||:|:|
Db 22 THHIIIEC 30

RESULT 24
US-11-096-568A-24717
; Sequence 24717, Application US/11096568A

; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; PRIOR FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24717
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(153)
; OTHER INFORMATION: Ceres Seq. ID no. 12450486
US-11-096-568A-24717

Query Match 69.4%; Score 34; DB 11; Length 153;
Best Local Similarity 57.1%; Pred. No. 1.4;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 3 HDIIIEC 9
|||:|:|
Db 126 HDIIIEC 132

RESULT 25
US-11-096-568A-21632
; Sequence 21632, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; PRIOR FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 21632
; LENGTH: 204
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(204)
; OTHER INFORMATION: Ceres Seq. ID no. 12405549
US-11-096-568A-21632

Query Match 69.4%; Score 34; DB 11; Length 204;
Best Local Similarity 62.5%; Pred. No. 1.9;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 2 HDIIIEC 9
|||:|:|
Db 104 HDIIIEC 111

RESULT 26
US-11-096-568A-24716
; Sequence 24716, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; PRIOR FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24716
; LENGTH: 210

```

; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(210)
; OTHER INFORMATION: Ceres Seq. ID no. 12450485
US-11-096-568A-24716
```

```
Query Match          69.4%; Score 34; DB 11; Length 210;
Best Local Similarity 57.1%; Pred. No. 20;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      3 HDIILLC 9
Db      183 HDIVLDC 189
```

```
RESULT 27
US-11-096-568A-5205
; Sequence 5205, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Theby
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 5205
; LENGTH: 264
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(264)
; OTHER INFORMATION: Ceres Seq. ID no. 14307434
US-11-096-568A-5205
```

```
Query Match          69.4%; Score 34; DB 11; Length 264;
Best Local Similarity 62.5%; Pred. No. 25;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
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```
Qy      1 TIDHILLC 8
Db      26 TIDHVLLE 33
```

```
RESULT 28
US-11-096-568A-21631
; Sequence 21631, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Theby
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 21631
; LENGTH: 310
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(310)
; OTHER INFORMATION: Ceres Seq. ID no. 12405548
US-11-096-568A-21631
```

```
Query Match          69.4%; Score 34; DB 11; Length 310;
Best Local Similarity 62.5%; Pred. No. 30;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy      2 IHDILLC 9
Db      210 IHDISIRC 217
```

```
RESULT 29
US-11-096-568A-21630
; Sequence 21630, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Theby
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 21630
; LENGTH: 311
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(311)
; OTHER INFORMATION: Ceres Seq. ID no. 12405547
US-11-096-568A-21630
```

```
Query Match          69.4%; Score 34; DB 11; Length 311;
Best Local Similarity 62.5%; Pred. No. 30;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy      2 IHDILLC 9
Db      211 IHDISIRC 218
```

```
RESULT 30
US-11-096-568A-5204
; Sequence 5204, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; TITLE OF INVENTION: Theby
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 5204
; LENGTH: 407
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc.feature
; LOCATION: (1)..(407)
; OTHER INFORMATION: Ceres Seq. ID no. 14307433
US-11-096-568A-5204
```

```
Query Match          69.4%; Score 34; DB 11; Length 407;
Best Local Similarity 62.5%; Pred. No. 40;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      1 TIDHILLC 8
Db      169 TIDHVLLE 176
```

```
RESULT 31
US-11-096-568A-5203
; Sequence 5203, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
```

```

; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: Inetdy
; CURRENT FILING DATE: 2005-11-02
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: 60/229,358
; PRIOR FILING DATE: 2000-04-12
; PRIOR APPLICATION NUMBER: 60/256,931
; PRIOR FILING DATE: 2000-12-21
; PRIOR APPLICATION NUMBER: 60/199,384
; PRIOR FILING DATE: 2000-04-25
; NUMBER OF SEQ ID NOS: 2267
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 110
; LENGTH: 157
; TYPE: PRT
; ORGANISM: Homo sapiens
; NAME/KEY: misc_feature
; LOCATION: (1)-(433)
; OTHER INFORMATION: Ceres Seq. ID no. 14307432
; US-11-096-568A-5203

Query Match      69.4%; Score 34; DB 11; Length 433;
Best Local Similarity 62.5%; Pred. No. 43;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy      1 THDIILE 8
      |||::||
Db      195 TLHDVLE 202

RESULT 32
US-10-858-730-70
; Sequence 70, Application US/10858730
; Publication No. US20050255568A1
; GENERAL INFORMATION:
; APPLICANT: Bailey, Richard B.
; APPLICANT: Blomquist, Paul
; APPLICANT: Doten, Reed
; APPLICANT: Diggers, Edward M.
; APPLICANT: Madden, Kevin T.
; APPLICANT: O'Leary, Jessica
; APPLICANT: O'Toole, George
; APPLICANT: Trueheart, Joshua
; APPLICANT: Walbridge, Michael J.
; APPLICANT: Yorgey, Peter S.
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR AMINO ACID
; FILE REFERENCE: 14184-030001
; CURRENT APPLICATION NUMBER: US/10/858,730
; CURRENT FILING DATE: 2004-06-01
; PRIOR APPLICATION NUMBER: US 60/475,000
; PRIOR FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US 60/551,860
; PRIOR FILING DATE: 2004-03-10
; NUMBER OF SEQ ID NOS: 364
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 70
; LENGTH: 1158
; TYPE: PRT
; ORGANISM: Thermobifida fusca
; US-10-858-730-70

Query Match      69.4%; Score 34; DB 9; Length 1158;
Best Local Similarity 71.4%; Pred. No. 1,2e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy      3 HDIILEC 9
      |||::||
Db      498 HDIIVDC 504

RESULT 33
US-11-264-096-110
; Sequence 110, Application US/11264096
; Publication No. US20060084794A1
; GENERAL INFORMATION:
; APPLICANT: Rosen et al.
```

```

; TITLE OF INVENTION: Albumin Fusion Proteins
; FILE REFERENCE: P5546D1
; CURRENT APPLICATION NUMBER: US/11/264,096
; CURRENT FILING DATE: 2005-11-02
; PRIOR APPLICATION NUMBER: 09/833,245
; PRIOR FILING DATE: 2001-04-12
; PRIOR APPLICATION NUMBER: 60/229,358
; PRIOR FILING DATE: 2000-04-12
; PRIOR APPLICATION NUMBER: 60/256,931
; PRIOR FILING DATE: 2000-12-21
; PRIOR APPLICATION NUMBER: 60/199,384
; PRIOR FILING DATE: 2000-04-25
; NUMBER OF SEQ ID NOS: 2267
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 110
; LENGTH: 157
; TYPE: PRT
; ORGANISM: Homo sapiens
; US-11-264-096-110

Query Match      67.3%; Score 33; DB 11; Length 157;
Best Local Similarity 71.4%; Pred. No. 23;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

Qy      2 IHDIILE 8
      |||::||
Db      67 IHDIIVLE 73

RESULT 34
US-11-172-740-1140
; Sequence 1140, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
; APPLICANT: Mascia, Peter
; APPLICANT: Alexandrov, Nickolai
; APPLICANT: Brover, Vyacheslav
; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
; FILE REFERENCE: 2750-1602PUS2
; CURRENT APPLICATION NUMBER: US/11/172,740
; CURRENT FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,800
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1140
; LENGTH: 281
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)-(281)
; OTHER INFORMATION: Ceres CLONE ID no. 463846
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Useful for increasing chlorophyll and photosynthetic cap
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Useful for making ornamental plants with modified flower
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Useful for making ornamental plants with modified leaves
```



```

; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making smaller plants
US-11-172-740-1140

```

```

Query Match
Best Local Similarity 37.5%; Score 33; DB 11; Length 281;
Matches 3; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

```

```

QY 2 IHDIILEC 9
DB 117 VHDVFIDC 124

```

```

RESULT 35
US-11-087-099-240
; Sequence 240, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 240
; LENGTH: 302
; TYPE: PRT
; ORGANISM: Glycine max
US-11-087-099-240

```

```

Query Match
Best Local Similarity 55.6%; Score 33; DB 11; Length 302;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

```

```

QY 1 TTHDIILEC 9
DB 192 TRHDIILRC 200

```

```

RESULT 36
US-11-087-099-5110
; Sequence 5110, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 5110
; LENGTH: 700
; TYPE: PRT
; ORGANISM: Saccharomyces cerevisiae
US-11-087-099-5110

```

```

Query Match
Best Local Similarity 44.4%; Score 33; DB 11; Length 700;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

```

```

QY 1 TTHDIILEC 9
DB 41 TYVEFYILEC 49

```

```

RESULT 37
US-11-079-463-8639
; Sequence 8639, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRAC
; FILE REFERENCE: PATH00-03DIV2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 8639
; LENGTH: 953
; TYPE: PRT
; ORGANISM: B.fragilis
US-11-079-463-8639

```

```

Query Match
Best Local Similarity 71.4%; Score 33; DB 11; Length 953;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 3 HDIILEC 9
DB 808 HEMILEC 814

```

```

RESULT 38
US-11-188-298-16409
; Sequence 16409, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16409
; LENGTH: 340
; TYPE: PRT
; ORGANISM: Prochlorococcus marinus subsp. pastoris str. CMP1378
US-11-188-298-16409

```

```

Query Match
Best Local Similarity 100.0%; Score 32; DB 11; Length 340;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

```

```

QY 4 DIILEC 9
DB 90 DIILEC 95

```

```

RESULT 39
US-11-188-298-646
; Sequence 646, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31

```

NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 646
LENGTH: 418
TYPE: PRT
ORGANISM: Burkholderia fungorum
US-11-188-298-646

Query Match 65.3%; Score 32; DB 11; Length 418;
Best Local Similarity 71.4%; Pred. No. 1e+02; 1; Indels 0; Gaps 0;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 3 HDIILC 9
Db 248 HDLILC 254

RESULT 40
US-10-505-928-102
Sequence 102, Application US/10505928
Publication No. US20060088532A1
GENERAL INFORMATION:
APPLICANT: Ludwig Institute for Cancer Research et al.
TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES
FILE REFERENCE: 28967/39178
CURRENT APPLICATION NUMBER: US/10/505,928
CURRENT FILING DATE: 2004-08-27
PRIOR APPLICATION NUMBER: US 60/363,019
PRIOR FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 866
SOFTWARE: PatentIn 3.2
SEQ ID NO 102
LENGTH: 780
TYPE: PRT
ORGANISM: Homo sapiens
US-10-505-928-102

Query Match 65.3%; Score 32; DB 8; Length 780;
Best Local Similarity 50.0%; Pred. No. 2e+02; 1; Indels 0; Gaps 0;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 2 HDIILC 9
Db 655 IHSVLDC 662

RESULT 41
US-10-055-877-139
Sequence 139, Application US/10055877
Publication No. US20050288241A1
GENERAL INFORMATION:
APPLICANT: Decistofaro, Marc
APPLICANT: Padigaru, Muralidhara
APPLICANT: Miller, Charles
APPLICANT: Tchervet, Veilizar
APPLICANT: Zhong, Mei
APPLICANT: Anderson, David
APPLICANT: Ballinger, Robert
APPLICANT: Gerlach, Valerie
APPLICANT: Spytek, Kimberly
APPLICANT: Raccelli, Luca
APPLICANT: Kekuda, Ramesh
APPLICANT: Guo, Xiaojia
APPLICANT: Zeehuusen, Bryan
APPLICANT: Andrew, David
APPLICANT: Mezes, Peter
APPLICANT: Paturajan, Meera
APPLICANT: Burgess, Catherine
APPLICANT: Eigen, Andrew
APPLICANT: Wolenc, Adam
APPLICANT: Baumgartner, Jason
APPLICANT: Shimkets, Richard
APPLICANT: Gusev, Vladimir
APPLICANT: Verneet, Corine

APPLICANT: Taupier Jr., Raymond
APPLICANT: Pena, Carol
APPLICANT: Shenoy, Suresh
APPLICANT: Li, Li
APPLICANT: Casman, Stacie
APPLICANT: Boldog, Ferenc
TITLE OF INVENTION: Novel Polypeptides and Nucleic Acids Encoded Thereby
FILE REFERENCE: 21402-251
CURRENT APPLICATION NUMBER: US/10/055,877
CURRENT FILING DATE: 2002-01-22
PRIOR APPLICATION NUMBER: 60/262,892
PRIOR FILING DATE: 2001-01-19
PRIOR APPLICATION NUMBER: 60/263,598
PRIOR FILING DATE: 2001-01-23
PRIOR APPLICATION NUMBER: 60/263,799
PRIOR FILING DATE: 2001-01-24
PRIOR APPLICATION NUMBER: 60/264,117
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,139
PRIOR FILING DATE: 2001-01-25
PRIOR APPLICATION NUMBER: 60/264,478
PRIOR FILING DATE: 2001-01-26
PRIOR APPLICATION NUMBER: 60/263,351
PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: 60/272,870
PRIOR FILING DATE: 2001-03-02
PRIOR APPLICATION NUMBER: 60/275,990
PRIOR FILING DATE: 2001-03-14
PRIOR APPLICATION NUMBER: 60/275,927
PRIOR FILING DATE: 2001-03-14
Remaining Prior Application data removed - See File Wrapper or PALM.

NUMBER OF SEQ ID NOS: 512
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 139
LENGTH: 1159
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-10-055-877-139

Query Match 65.3%; Score 32; DB 9; Length 1159;
Best Local Similarity 50.0%; Pred. No. 3e+02; 1; Indels 0; Gaps 0;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 2 HDIILC 9
Db 1044 HDVLOQC 1051

RESULT 42
US-10-467-657-5966
Sequence 5966, Application US/10467657
Publication No. US20050260581A1
GENERAL INFORMATION:
APPLICANT: CHIRON SpA
APPLICANT: FONTANA Maria Rita
APPLICANT: PIZZA Mariagrazia
APPLICANT: MASTIGNI Vega
APPLICANT: MONACT Elisabetta
TITLE OF INVENTION: GONOCOCCAL PROTEINS AND NUCLEIC ACIDS
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/10/467,657
CURRENT FILING DATE: 2003-08-11
PRIOR APPLICATION NUMBER: GB-0103424.8
PRIOR FILING DATE: 2001-02-12
NUMBER OF SEQ ID NOS: 9218
SOFTWARE: SeqWin99, version 1.04
SEQ ID NO 5966
LENGTH: 80
TYPE: PRT
ORGANISM: Neisseria gonorrhoeae
US-10-467-657-5966

Query Match 63.3%; Score 31; DB 9; Length 80;

Best Local Similarity 55.6%; Pred. No. 28;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy 1 TTHDIIIEC 9
|:|||||
Db 37 TLNQIILDC 45

RESULT 43

US-10-530-253-18
; Sequence 18, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 35
US-10-530-253-18

Query Match 63.3%; Score 31; DB 9; Length 149;
Best Local Similarity 55.6%; Pred. No. 55;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 TTHDIIIEC 9
|:|||||
Db 22 SIHEICLNC 30

RESULT 44

US-10-491-468-22
; Sequence 22, Application US/10491468
; Publication No. US20060051836A1
; GENERAL INFORMATION:
; APPLICANT: INCYTE CORPORATION; TANG, Y. Tom;
; APPLICANT: FORSYTHE, Ian J.; EMERLING, Brooke M.;
; APPLICANT: HAPALIA, April J.A.; YUE, Henry;
; APPLICANT: XU, Yuming; GIERZEN, Kimberly J.;
; APPLICANT: CHAWLA, Narinder K.; BAUGHN, Mariah R.;
; APPLICANT: MARQUIS, Joseph P.; BECHA, Shanya D.;
; APPLICANT: KABLE, Amy E.; LAU, Preeti G.;
; APPLICANT: RICHARDSON, Thomas W.; LEE, Soo Y.;
; APPLICANT: LEE, Ernestine A.; TRAN, Bao;
; APPLICANT: WARREN, Bridget A.; LU, Dyung Anna M.;
; APPLICANT: GURURAJAN, Rajasopali; SPRAGUE, William W.;
; APPLICANT: BLAKE, Julie J.; THANGAVELU, Kavitha;
; APPLICANT: SWARNAKAR, Anita; GORVAD, Ann E.;
; APPLICANT: GRIFPIN, Jennifer A.; LINDQUIST, Erika A.;
; APPLICANT: ELLIOTT, Vicki S.; ISON, Craig H.;
; APPLICANT: RAMKUNAR, Jeyalaksmi
; TITLE OF INVENTION: MOLECULES FOR DISEASE DETECTION AND TREATMENT
; FILE REFERENCE: PF-1232 USN
; CURRENT APPLICATION NUMBER: US/10/491,468
; CURRENT FILING DATE: 2004-03-31
; PRIOR APPLICATION NUMBER: PCT/US02/32852
; PRIOR FILING DATE: 2002-10-10
; PRIOR APPLICATION NUMBER: US 60/328,944
; PRIOR FILING DATE: 2001-10-12
; PRIOR APPLICATION NUMBER: US 60/345,304
; PRIOR FILING DATE: 2001-10-26

; PRIOR APPLICATION NUMBER: US 60/343,880
; PRIOR FILING DATE: 2001-11-02
; PRIOR APPLICATION NUMBER: US 60/345,143
; PRIOR FILING DATE: 2001-11-09
; PRIOR APPLICATION NUMBER: US 60/332,430
; PRIOR FILING DATE: 2001-11-16
; NUMBER OF SEQ ID NOS: 96
; SOFTWARE: PERL Program
; SEQ ID NO 22
; LENGTH: 234
; TYPE: PRT
; ORGANISM: Homo sapiens
; NAME/KEY: misc.feature
; OTHER INFORMATION: Incyte ID No: 5281724CD1
US-10-491-468-22

Query Match 63.3%; Score 31; DB 9; Length 234;
Best Local Similarity 71.4%; Pred. No. 88;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 3 HDIIIEC 9
|:|||||
Db 87 HPIILIC 93

RESULT 45

US-10-878-556A-177
; Sequence 177, Application US/10878556A
; Publication No. US20050266399A1
; GENERAL INFORMATION:
; APPLICANT: Hoffmann La-Roche Inc.
; TITLE OF INVENTION: HCV regulated protein expression
; FILE REFERENCE: 21762
; CURRENT APPLICATION NUMBER: US/10/878,556A
; CURRENT FILING DATE: 2004-06-28
; NUMBER OF SEQ ID NOS: 199
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 177
; LENGTH: 254
; TYPE: PRT
; ORGANISM: Homo sapiens
; PUBLICATION INFORMATION:
; DATABASE ACCESSION NUMBER: sw_hum/peet3_human
; DATABASE ENTRY DATE: 2001-10-16
US-10-878-556A-177

Query Match 63.3%; Score 31; DB 9; Length 254;
Best Local Similarity 62.5%; Pred. No. 96;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 1 TTHDIIIEC 8
|:|||||
Db 230 TLHDMILK 237

RESULT 46

US-11-096-568A-4430
; Sequence 4430, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 4430
; LENGTH: 264
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:

```
; NAME/KEY: misc_feature
; LOCATION: (1) (264)
; OTHER INFORMATION: Ceres Seq. ID no. 15219381
US-11-096-568A-4430
```

```
Query Match      63.3%; Score 31; DB 11; Length 264;
Best Local Similarity 57.1%; Pred. No. 1e+02;
Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      3 HDIILEC 9
      ||::||
Db      151 HDILIRC 157
```

```
RESULT 47
US-11-096-568A-24104
; Sequence 24104, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2/50-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; PRIOR FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24104
; LENGTH: 267
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1) (267)
; OTHER INFORMATION: Ceres Seq. ID no. 12418821
US-11-096-568A-24104
```

```
Query Match      63.3%; Score 31; DB 11; Length 267;
Best Local Similarity 57.1%; Pred. No. 1e+02;
Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      3 HDIILEC 9
      ||::||
Db      220 HDCTVEC 226
```

```
RESULT 48
US-10-505-928-708
; Sequence 708, Application US/10505928
; Publication No. US20060088532A1
; GENERAL INFORMATION:
; APPLICANT: Ludwig Institute for Cancer Research et al.
; TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES
; FILE REFERENCE: 28967/39178
; CURRENT APPLICATION NUMBER: US/10/505,928
; PRIOR FILING DATE: 2004-08-27
; PRIOR APPLICATION NUMBER: US 60/363,019
; PRIOR FILING DATE: 2002-03-07
; NUMBER OF SEQ ID NOS: 866
; SOFTWARE: PatentIn 3.2
; SEQ ID NO 708
; LENGTH: 275
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-505-928-708
```

```
Query Match      63.3%; Score 31; DB 8; Length 275;
Best Local Similarity 57.1%; Pred. No. 1e+02;
Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      3 HDIILEC 9
      ||::||
Db      85 HDTVLQC 91
```

```
RESULT 49
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; Sequence 710, Application US/10505928
; Publication No. US20060088532A1
; GENERAL INFORMATION:
; APPLICANT: Ludwig Institute for Cancer Research et al.
; TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES
; FILE REFERENCE: 28967/39178
; CURRENT APPLICATION NUMBER: US/10/505,928
; PRIOR FILING DATE: 2004-08-27
; PRIOR APPLICATION NUMBER: US 60/363,019
; PRIOR FILING DATE: 2002-03-07
; NUMBER OF SEQ ID NOS: 866
; SOFTWARE: PatentIn 3.2
; SEQ ID NO 710
; LENGTH: 275
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-505-928-710
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Best Local Similarity 57.1%; Pred. No. 1e+02;
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Db      85 HDTVLQC 91
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RESULT 50
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; Sequence 64, Application US/10493909
; Publication No. US20060015969A1
; GENERAL INFORMATION:
; APPLICANT: LARRICK, JAMES W.
; TITLE OF INVENTION: NOVEL IMMUNOADHESINS FOR TREATING AND PREVENTING TOXICITY
; FILE REFERENCE: 41514-20004.01
; CURRENT APPLICATION NUMBER: US/10/493,909
; PRIOR FILING DATE: 2004-04-26
; PRIOR APPLICATION NUMBER: PCT/US01/13932
; PRIOR FILING DATE: 2001-04-28
; PRIOR APPLICATION NUMBER: 60/200,298
; PRIOR FILING DATE: 2000-04-28
; NUMBER OF SEQ ID NOS: 101
; SOFTWARE: PatentIn Ver. 2.1
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; LENGTH: 275
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-493-909-64
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Query Match      63.3%; Score 31; DB 9; Length 275;
Best Local Similarity 57.1%; Pred. No. 1e+02;
Matches 4; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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Db      85 HDTVLQC 91
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Search completed: May 5, 2006, 08:18:50
Job time : 9.4 secs
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GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 04:01:20 ; Search time 20.7 Seconds
(without alignments)
35.946 Million cell updates/sec

Title:	US-08-170-344-8
Perfect score:	48
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Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

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Searched:      572060 seqs, 82675679 residues
Total number of hits satisfying chosen parameters: 572060
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Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed and is derived by analysts of the total score distribution.

SUMMARIES

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4	48	100.0	10	4	PCT-US95-02121-72	Sequence 72, Appl
5	48	100.0	30	2	US-09-980-523A-4	Sequence 4, Appl
6	48	100.0	59	2	US-09-390-027-6	Sequence 6, Appl
7	48	100.0	151	2	US-09-701-080C-18	Sequence 18, Appl
8	48	100.0	158	2	US-09-980-523A-2	Sequence 2, Appl
9	48	100.0	162	1	US-08-316-229B-3	Sequence 3, Appl
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168	30	62.5	1436	1	US-08-769-399-2	Sequence 2, Appli	241	29	60.4	434	2	US-09-811-469-5	Sequence 5, Appli
169	30	62.5	1436	1	US-08-991-258A-2	Sequence 2, Appli	242	29	60.4	476	2	US-10-370-659-4	Sequence 4, Appli
170	30	62.5	1436	1	US-08-991-953A-2	Sequence 2, Appli	243	29	60.4	476	2	US-09-252-991A-22879	Sequence 22879, A
171	30	62.5	1439	1	US-08-449-644-2	Sequence 2, Appli	244	29	60.4	485	2	US-08-426-630-43	Sequence 43, Appli
172	30	62.5	1439	1	US-08-087-244A-2	Sequence 2, Appli	245	29	60.4	492	2	US-09-248-796A-16231	Sequence 16231, A
173	30	62.5	1452	1	US-08-652-971-4	Sequence 4, Appli	246	29	60.4	492	2		

247	29	60.4	530	2	US-09-444-711A-4	Sequence 4, Appl1	320	28	58.3	153	2	US-09-270-767-33188	Sequence 33188, A
248	29	60.4	531	2	US-09-134-000C-5332	Sequence 5322, Ap	321	28	58.3	156	2	US-09-270-767-48717	Sequence 48717, A
249	29	60.4	532	2	US-09-902-540-9698	Sequence 9698, A	322	28	58.3	159	2	US-09-270-767-32558	Sequence 32558, A
250	29	60.4	533	1	US-07-820-011A-2	Sequence 2, Appl1	323	28	58.3	159	2	US-09-270-767-47775	Sequence 47775, A
251	29	60.4	533	2	US-09-470-881-3	Sequence 2, Appl1	324	28	58.3	162	2	US-09-270-767-31615	Sequence 31813, A
252	29	60.4	533	4	PCT-US93-00445-2	Sequence 2, Appl1	325	28	58.3	162	2	US-09-270-767-47032	Sequence 47032, A
253	29	60.4	536	1	US-07-820-011A-4	Sequence 4, Appl1	326	28	58.3	170	2	US-09-489-039A-7747	Sequence 7747, Ap
254	29	60.4	536	2	US-08-426-509A-13	Sequence 11, Appl	327	28	58.3	174	2	US-09-248-796A-20708	Sequence 20708, A
255	29	60.4	536	2	US-08-232-545-13	Sequence 11, Appl	328	28	58.3	172	2	US-09-248-796A-20508	Sequence 20508, A
256	29	60.4	536	2	US-09-444-711A-2	Sequence 2, Appl1	329	28	58.3	184	2	US-09-124-864-56	Sequence 56, Appl
257	29	60.4	536	2	US-09-929-266-10	Sequence 10, Appl	330	28	58.3	198	2	US-09-252-991A-32955	Sequence 32955, A
258	29	60.4	536	2	US-09-977-261-13	Sequence 11, Appl	331	28	58.3	204	2	US-09-134-001C-2975	Sequence 2975, Ap
259	29	60.4	536	4	PCT-US93-00445-4	Sequence 4, Appl1	332	28	58.3	206	2	US-09-949-016-10713	Sequence 10713, A
260	29	60.4	536	4	PCT-US93-00445-4	Sequence 11, Appl	333	28	58.3	219	2	US-09-134-001C-2975	Sequence 2975, Ap
261	29	60.4	537	2	US-09-949-016-10282	Sequence 10282, A	334	28	58.3	219	2	US-09-134-001C-2975	Sequence 2975, Ap
262	29	60.4	641	2	US-09-919-039-73	Sequence 73, Appl	335	28	58.3	219	2	US-09-134-001C-2975	Sequence 2975, Ap
263	29	60.4	646	1	US-08-441-139-11	Sequence 11, Appl	336	28	58.3	219	2	US-09-134-001C-2975	Sequence 2975, Ap
264	29	60.4	646	2	US-09-919-039-11	Sequence 11, Appl	337	28	58.3	219	2	US-09-134-001C-2975	Sequence 2975, Ap
265	29	60.4	681	2	US-10-160-748-2	Sequence 2, Appl1	338	28	58.3	243	2	US-09-328-352-6052	Sequence 6052, Ap
266	29	60.4	748	2	US-09-949-016-10387	Sequence 10387, A	339	28	58.3	245	2	US-08-015-985-8	Sequence 8, Appl1
267	29	60.4	777	2	US-09-811-469-2	Sequence 2, Appl1	340	28	58.3	245	2	US-09-280-597-8	Sequence 8, Appl1
268	29	60.4	777	2	US-10-370-659-2	Sequence 2, Appl1	341	28	58.3	260	1	US-08-685-992-4	Sequence 25, Appl
269	29	60.4	780	1	US-08-887-798-2	Sequence 2, Appl1	342	28	58.3	260	1	US-08-685-992-25	Sequence 25, Appl
270	29	60.4	786	2	US-09-949-016-10898	Sequence 10898, A	343	28	58.3	260	1	US-09-144-925-25	Sequence 25, Appl
271	29	60.4	841	2	US-10-332-795-11	Sequence 11, Appl	344	28	58.3	261	1	US-09-144-925-3	Sequence 3, Appl1
272	29	60.4	883	2	US-09-248-796A-14418	Sequence 14418, A	345	28	58.3	261	1	US-08-685-992-3	Sequence 9, Appl1
273	29	60.4	890	2	US-09-513-783A-174	Sequence 174, App	346	28	58.3	262	1	US-08-241-766-9	Sequence 24, Appl
274	29	60.4	890	2	US-10-100-957A-174	Sequence 174, App	347	28	58.3	262	1	US-10-089-019-24	Sequence 6343, Ap
275	29	60.4	894	2	US-09-949-016-10605	Sequence 10605, A	348	28	58.3	271	2	US-09-328-352-6343	Sequence 108, Ap
276	29	60.4	997	2	US-09-198-452A-602	Sequence 602, App	349	28	58.3	272	2	US-09-107-532A-4013	Sequence 4013, Ap
277	29	60.4	997	2	US-09-900-920-60	Sequence 60, App	350	28	58.3	275	2	US-09-107-532A-4013	Sequence 4013, Ap
278	29	60.4	1125	2	US-09-919-891-2	Sequence 339, App	351	28	58.3	275	2	US-09-107-532A-4013	Sequence 4013, Ap
279	29	60.4	1358	2	US-09-538-092-339	Sequence 339, App	352	28	58.3	275	2	US-09-107-532A-4013	Sequence 4013, Ap
280	29	60.4	1405	2	US-09-438-185A-566	Sequence 566, App	353	28	58.3	280	2	US-10-003-690-8	Sequence 8, Appl1
281	29	60.4	1474	2	US-09-677-046A-4	Sequence 4, Appl1	354	28	58.3	280	2	US-09-515-806-15	Sequence 15, Appl
282	29	60.4	1503	2	US-09-677-046A-6	Sequence 6, Appl1	355	28	58.3	286	2	US-09-252-991A-17714	Sequence 17714, A
283	29	60.4	1509	2	US-09-677-046A-2	Sequence 2, Appl1	356	28	58.3	287	2	US-09-134-001C-4454	Sequence 4454, Ap
284	29	60.4	2237	1	US-08-455-543A-48	Sequence 48, Appl	357	28	58.3	291	2	US-09-107-532A-5063	Sequence 5063, Ap
285	29	60.4	2237	1	US-08-223-305C-48	Sequence 48, Appl	358	28	58.3	308	2	US-10-014-269-31	Sequence 31, Appl
286	29	60.4	2237	2	US-09-268-163-8	Sequence 8, Appl1	359	28	58.3	308	2	US-10-002-974-31	Sequence 31, Appl
287	29	60.4	2262	2	US-09-949-016-4849	Sequence 849, Ap	360	28	58.3	312	2	US-09-270-767-44669	Sequence 44669, A
288	29	60.4	2336	2	US-09-268-163-10	Sequence 10, Appl	361	28	58.3	313	2	US-08-480-640A-192	Sequence 192, App
289	29	60.4	2337	2	US-08-713-118-2	Sequence 2, Appl1	362	28	58.3	313	2	US-08-686-968C-192	Sequence 192, App
290	29	60.4	2337	2	US-09-452-007-2	Sequence 2, Appl1	363	28	58.3	313	2	US-08-375-992A-192	Sequence 192, App
291	29	60.4	2339	1	US-08-455-543A-47	Sequence 47, Appl1	364	28	58.3	313	2	US-08-375-992A-192	Sequence 192, App
292	29	60.4	2339	1	US-08-223-305C-47	Sequence 47, Appl1	365	28	58.3	314	2	US-09-902-540-16180	Sequence 16180, A
293	29	60.4	2339	2	US-09-268-163-6	Sequence 6, Appl1	366	28	58.3	317	2	US-09-248-796A-17349	Sequence 17349, A
294	29	60.4	2343	2	US-09-268-163-4	Sequence 4, Appl1	367	28	58.3	326	1	US-09-055-097-6	Sequence 6, Appl1
295	29	60.4	3066	2	US-08-952-127-12	Sequence 12, Appl	368	28	58.3	326	1	US-09-373-902-6	Sequence 11, Appl
296	29	60.4	374	2	US-09-134-001C-2939	Sequence 2939, Ap	369	28	58.3	326	2	US-09-831-630-11	Sequence 25, Appl
297	28.5	59.4	574	1	US-08-907-166-12	Sequence 12, Appl	370	28	58.3	331	2	US-08-556-419-25	Sequence 23980, A
298	28.5	59.4	574	2	US-09-391-340-12	Sequence 20, Appl	371	28	58.3	331	2	US-09-252-991A-23980	Sequence 23980, A
299	28	58.3	29	2	US-08-556-419-20	Sequence 20, Appl	372	28	58.3	344	2	US-09-107-532A-3773	Sequence 3773, Ap
300	28	58.3	29	2	US-09-962-766-1365	Sequence 1365, Ap	373	28	58.3	344	2	US-09-107-532A-3773	Sequence 3773, Ap
301	28	58.3	33	2	US-09-270-767-34106	Sequence 34106, Ap	374	28	58.3	354	2	US-09-810-836B-4	Sequence 4366, Ap
302	28	58.3	33	2	US-09-270-767-49323	Sequence 49323, A	375	28	58.3	361	2	US-09-134-000C-4306	Sequence 10712, Ap
307	28	58.3	88	2	US-09-270-767-36177	Sequence 36177, A	376	28	58.3	362	2	US-09-538-092-1072	Sequence 10714, A
308	28	58.3	88	2	US-09-270-767-51394	Sequence 51394, A	377	28	58.3	363	2	US-09-949-016-11214	Sequence 14, Appl
309	28	58.3	88	2	US-09-248-796A-25949	Sequence 25949, A	378	28	58.3	363	2	US-09-949-016-11214	Sequence 14, Appl
310	28	58.3	88	2	US-08-928-383B-21	Sequence 21, Appl	379	28	58.3	371	1	US-08-673-789-11	Sequence 2, Appl1
311	28	58.3	96	2	US-08-478-208-4	Sequence 4, Appl1	380	28	58.3	371	1	US-08-673-789-11	Sequence 2, Appl1
312	28	58.3	96	2	US-08-921-195-4	Sequence 4, Appl1	381	28	58.3	389	2	US-08-488-217A-114	Sequence 114, App
313	28	58.3	106	2	US-09-152-060-90	Sequence 90, Appl	382	28	58.3	389	2	US-08-295-802-114	Sequence 114, App
314	28	58.3	106	2	US-09-852-797-90	Sequence 90, Appl	383	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App
315	28	58.3	106	2	US-09-853-161-90	Sequence 90, Appl	384	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App
316	28	58.3	106	2	US-10-056-993-90	Sequence 90, Appl	385	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App
317	28	58.3	126	2	US-08-556-419-24	Sequence 24, Appl	386	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App
318	28	58.3	136	2	US-09-270-767-55283	Sequence 55283, A	387	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App
319	28	58.3	145	2	US-09-270-767-48404	Sequence 48404, A	388	28	58.3	389	2	US-08-472-619H-114	Sequence 114, App

393	28	58.3	396	2	US-09-248-796A-18977	Sequence 18977, A	466	28	58.3	661	1	US-08-795-868-14	Sequence 14, Appl
394	28	58.3	397	2	US-09-949-016-11445	Sequence 11445, A	467	28	58.3	661	2	US-09-303-069-14	Sequence 14, Appl
395	28	58.3	398	1	US-08-091-519-2	Sequence 2, Appl	468	28	58.3	661	2	US-09-134-250-14	Sequence 14, Appl
396	28	58.3	398	1	US-08-442-043A-2	Sequence 2, Appl	469	28	58.3	671	2	US-09-949-016-6441	Sequence 6441, Ap
397	28	58.3	398	2	US-09-173-151A-26	Sequence 26, Appl	470	28	58.3	674	2	US-09-949-016-7034	Sequence 7034, Ap
398	28	58.3	398	2	US-09-461-908-2	Sequence 2, Appl	471	28	58.3	682	1	US-08-911-445-3	Sequence 3, Appl
399	28	58.3	398	2	US-08-441-893A-8	Sequence 2, Appl	472	28	58.3	682	2	US-09-182-983-3	Sequence 3, Appl
400	28	58.3	398	2	US-08-406-824A-8	Sequence 8, Appl	473	28	58.3	682	2	US-09-771-838A-3	Sequence 3, Appl
401	28	58.3	398	2	US-09-921-667-8	Sequence 8, Appl	474	28	58.3	682	2	US-09-949-016-6591	Sequence 6591, Ap
402	28	58.3	398	4	PCT-US91-03478-2	Sequence 2, Appl	475	28	58.3	682	2	US-09-949-016-6652	Sequence 6652, Ap
403	28	58.3	409	1	US-08-190-802A-51	Sequence 51, Appl	476	28	58.3	699	2	US-10-144-198-40	Sequence 40, Appl
404	28	58.3	409	1	US-08-283-917-3	Sequence 3, Appl	477	28	58.3	710	2	US-09-949-016-6808	Sequence 6808, Ap
405	28	58.3	409	1	US-08-961-716-3	Sequence 3, Appl	478	28	58.3	710	2	US-10-144-198-39	Sequence 39, Appl
406	28	58.3	409	2	US-08-477-346-51	Sequence 51, Appl	479	28	58.3	733	2	US-09-964-992A-3	Sequence 2, Appl
407	28	58.3	409	2	US-08-473-089-51	Sequence 51, Appl	480	28	58.3	733	2	US-09-270-767-43994	Sequence 43994, A
408	28	58.3	409	2	US-08-487-072A-51	Sequence 51, Appl	481	28	58.3	736	2	US-09-270-767-43721	Sequence 43721, A
409	28	58.3	409	2	US-09-538-092-1119	Sequence 1119, Ap	482	28	58.3	736	2	US-09-949-016-6807	Sequence 6807, Ap
410	28	58.3	410	1	US-08-283-917-9	Sequence 9, Appl	483	28	58.3	736	2	US-10-144-198-38	Sequence 38, Appl
411	28	58.3	410	1	US-08-961-716-9	Sequence 9, Appl	484	28	58.3	738	2	US-08-478-208-32	Sequence 32, Appl
412	28	58.3	410	2	US-09-252-991A-33110	Sequence 33110, A	485	28	58.3	738	2	US-09-336-536-73	Sequence 73, Appl
413	28	58.3	414	2	US-09-107-532A-5771	Sequence 5771, Ap	486	28	58.3	752	2	US-10-144-198-22	Sequence 22, Appl
414	28	58.3	418	2	US-09-949-016-9458	Sequence 9458, Ap	487	28	58.3	763	2	US-10-144-198-20	Sequence 20, Appl
415	28	58.3	426	2	US-09-248-796A-17266	Sequence 17266, A	488	28	58.3	773	1	US-08-019-870-1	Sequence 1, Appl
416	28	58.3	430	2	US-08-686-968C-3	Sequence 3, Appl	489	28	58.3	774	1	US-08-019-870-6	Sequence 6, Appl
417	28	58.3	432	2	US-09-326-203A-25	Sequence 25, Appl	490	28	58.3	774	1	US-07-747-901A-3	Sequence 3, Appl
418	28	58.3	435	2	US-09-491-577-54	Sequence 54, Appl	491	28	58.3	774	1	US-07-935-312-3	Sequence 3, Appl
419	28	58.3	439	2	US-08-975-762-50	Sequence 60, Appl	492	28	58.3	774	1	US-08-019-870-5	Sequence 5, Appl
420	28	58.3	439	2	US-09-295-028-60	Sequence 60, Appl	493	28	58.3	774	1	US-08-019-870-8	Sequence 8, Appl
421	28	58.3	439	2	US-09-106-582-60	Sequence 60, Appl	494	28	58.3	774	1	US-08-019-870-11	Sequence 11, Appl
422	28	58.3	439	2	US-09-159-469-60	Sequence 60, Appl	495	28	58.3	774	1	US-07-731-157A-7	Sequence 7, Appl
423	28	58.3	439	2	US-09-693-542-60	Sequence 60, Appl	496	28	58.3	774	1	US-08-314-309A-21	Sequence 21, Appl
424	28	58.3	440	1	US-08-061-062A-6	Sequence 6, Appl	497	28	58.3	774	1	US-08-633-760-44	Sequence 44, Appl
425	28	58.3	440	1	US-08-061-062A-8	Sequence 8, Appl	498	28	58.3	774	1	US-08-633-760-46	Sequence 46, Appl
426	28	58.3	440	2	US-08-536-150-6	Sequence 6, Appl	499	28	58.3	774	1	US-08-633-760-48	Sequence 48, Appl
427	28	58.3	440	2	US-08-536-150-8	Sequence 8, Appl	500	28	58.3	774	1	US-08-633-760-50	Sequence 50, Appl
428	28	58.3	441	1	US-09-492-709A-272	Sequence 272, Appl	501	28	58.3	774	1	US-08-633-760-52	Sequence 52, Appl
429	28	58.3	445	2	US-09-047-026A-6	Sequence 6, Appl	502	28	58.3	774	1	US-08-541-780-7	Sequence 7, Appl
430	28	58.3	449	2	US-09-830-220A-269	Sequence 269, Appl	503	28	58.3	779	2	US-10-144-198-18	Sequence 18, Appl
431	28	58.3	463	1	US-09-248-796A-14957	Sequence 14957, A	504	28	58.3	802	2	US-09-134-001C-3840	Sequence 3840, Ap
432	28	58.3	466	1	US-08-785-066-2	Sequence 2, Appl	505	28	58.3	817	2	US-09-252-991A-20757	Sequence 20757, A
433	28	58.3	466	2	US-09-007-355-2	Sequence 2, Appl	506	28	58.3	849	1	US-08-162-809-6	Sequence 6, Appl
434	28	58.3	466	2	US-08-913-489-2	Sequence 2, Appl	507	28	58.3	849	1	US-08-162-809-9	Sequence 9, Appl
435	28	58.3	474	2	US-09-538-092-559	Sequence 559, Appl	508	28	58.3	869	2	US-08-673-789-10	Sequence 10, Appl
436	28	58.3	475	2	US-09-328-352-4223	Sequence 4223, Ap	509	28	58.3	869	2	US-10-104-047-2532	Sequence 2532, Ap
437	28	58.3	479	2	US-09-134-001C-4128	Sequence 4128, Ap	510	28	58.3	883	2	US-09-949-016-9129	Sequence 9129, Ap
438	28	58.3	492	2	US-09-489-039A-11838	Sequence 11838, A	511	28	58.3	915	2	US-10-282-162-46	Sequence 46, Appl
439	28	58.3	499	2	US-08-426-509A-19	Sequence 19, Appl	512	28	58.3	915	2	US-10-282-162-52	Sequence 52, Appl
440	28	58.3	499	2	US-08-232-545-19	Sequence 19, Appl	513	28	58.3	917	2	US-10-282-162-48	Sequence 48, Appl
441	28	58.3	499	2	US-09-977-261-19	Sequence 19, Appl	514	28	58.3	917	2	US-10-282-162-50	Sequence 50, Appl
442	28	58.3	499	4	PCT-US95-05008-19	Sequence 19, Appl	515	28	58.3	917	2	US-10-282-162-54	Sequence 54, Appl
443	28	58.3	509	2	US-09-902-540-11469	Sequence 11469, A	516	28	58.3	917	2	US-10-282-162-56	Sequence 56, Appl
444	28	58.3	512	2	US-09-390-234-24	Sequence 24, Appl	517	28	58.3	934	2	US-09-252-991A-25635	Sequence 25635, A
445	28	58.3	512	2	US-09-603-311-24	Sequence 24, Appl	518	28	58.3	945	2	US-09-949-016-8172	Sequence 8172, Ap
446	28	58.3	512	2	US-09-570-856B-30	Sequence 30, Appl	519	28	58.3	951	1	US-08-162-809-2	Sequence 2, Appl
447	28	58.3	513	2	US-09-732-025-2	Sequence 2, Appl	520	28	58.3	956	2	US-09-949-016-9128	Sequence 9128, Ap
448	28	58.3	513	2	US-10-153-917-2	Sequence 2, Appl	521	28	58.3	970	1	US-08-673-789-7	Sequence 7, Appl
449	28	58.3	527	2	US-09-739-455-2	Sequence 2, Appl	522	28	58.3	970	1	US-08-449-645A-11	Sequence 11, Appl
450	28	58.3	527	2	US-10-153-919-2	Sequence 2, Appl	523	28	58.3	970	1	US-08-702-367A-11	Sequence 11, Appl
451	28	58.3	530	2	US-09-248-796A-17374	Sequence 17374, A	524	28	58.3	974	4	PCT-US95-04681-11	Sequence 11, Appl
452	28	58.3	541	2	US-09-641-690A-2	Sequence 2, Appl	525	28	58.3	984	1	US-08-162-809-10	Sequence 10, Appl
453	28	58.3	581	2	US-10-104-047-3540	Sequence 3540, Ap	526	28	58.3	984	1	US-08-673-789-6	Sequence 6, Appl
454	28	58.3	599	2	US-08-556-419-22	Sequence 22, Appl	527	28	58.3	984	2	US-09-949-016-6502	Sequence 6502, Ap
455	28	58.3	601	1	US-08-795-868-16	Sequence 16, Appl	528	28	58.3	986	1	US-08-673-789-3	Sequence 3, Appl
456	28	58.3	602	1	US-09-303-069-16	Sequence 16, Appl	529	28	58.3	986	1	US-08-449-645A-15	Sequence 15, Appl
457	28	58.3	602	2	US-09-134-250-16	Sequence 16, Appl	530	28	58.3	986	1	US-08-702-367A-15	Sequence 15, Appl
458	28	58.3	608	2	US-09-270-767-43297	Sequence 43297, A	531	28	58.3	986	4	PCT-US95-04681-15	Sequence 15, Appl
459	28	58.3	615	1	US-08-911-445-2	Sequence 2, Appl	532	28	58.3	987	1	US-08-436-044-6	Sequence 6, Appl
460	28	58.3	615	2	US-09-182-983-2	Sequence 2, Appl	533	28	58.3	987	1	US-08-436-054-6	Sequence 6, Appl
461	28	58.3	615	2	US-09-771-838A-2	Sequence 2, Appl	534	28	58.3	987	4	PCT-US95-08812-6	Sequence 6, Appl
462	28	58.3	623	2	US-09-710-279-1068	Sequence 1068, Ap	535	28	58.3	988	1	US-08-162-809-14	Sequence 14, Appl
463	28	58.3	629	2	US-08-556-419-23	Sequence 23, Appl	536	28	58.3	990	1	US-09-949-016-7235	Sequence 7235, Ap
464	28	58.3	643	2	US-08-797-358B-3	Sequence 3, Appl	537	28	58.3	991	2	US-09-689-486-52	Sequence 52, Appl
465	28	58.3	643	2	US-09-919-039-204	Sequence 204, Appl	538	28	58.3	992	2	US-09-689-486-5	Sequence 5, Appl

539	28	58.3	992	2	US-09-689-486-53	Sequence 53, Appl	612	28	58.3	2314	2	US-09-816-703A-2	Sequence 2, Appl1
540	28	58.3	993	1	US-08-348-143-1	Sequence 1, Appl1	613	28	58.3	2509	1	US-08-149-097D-35	Sequence 35, Appl
541	28	58.3	993	1	US-08-571-785-1	Sequence 1, Appl1	614	28	58.3	2546	2	US-09-949-016-9500	Sequence 9500, Ap
542	28	58.3	993	2	US-08-368-776A-11	Sequence 11, Appl	615	28	58.3	2546	2	US-09-949-016-9501	Sequence 9501, Ap
543	28	58.3	993	2	US-09-192-435-1	Sequence 1, Appl1	616	28	58.3	2842	1	US-07-741-940-7	Sequence 7, Appl1
544	28	58.3	993	2	US-09-558-340-1	Sequence 1, Appl1	617	28	58.3	2842	1	US-08-289-548A-7	Sequence 7, Appl1
545	28	58.3	994	2	US-08-542-635-2	Sequence 2, Appl1	618	28	58.3	2842	1	US-08-452-654-7	Sequence 7, Appl1
546	28	58.3	994	2	US-08-368-776A-12	Sequence 12, Appl	619	28	58.3	2842	2	US-08-449-731-7	Sequence 7, Appl1
547	28	58.3	995	1	US-08-162-809-18	Sequence 18, Appl	620	28	58.3	2843	1	US-07-741-940-2	Sequence 2, Appl1
548	28	58.3	995	1	US-08-673-789-5	Sequence 5, Appl1	621	28	58.3	2843	1	US-08-289-548A-2	Sequence 2, Appl1
549	28	58.3	997	2	US-09-949-016-7171	Sequence 7171, Ap	622	28	58.3	2843	1	US-08-452-654-2	Sequence 2, Appl1
550	28	58.3	998	1	US-08-449-645A-17	Sequence 17, Appl	623	28	58.3	2843	1	US-08-452-655B-2	Sequence 2, Appl1
551	28	58.3	998	1	US-08-449-645A-20	Sequence 20, Appl	624	28	58.3	2843	1	US-08-452-655B-7	Sequence 7, Appl1
552	28	58.3	998	1	US-08-702-367A-17	Sequence 17, Appl	625	28	58.3	2843	1	US-08-370-235A-2	Sequence 2, Appl1
553	28	58.3	998	1	US-08-702-367A-20	Sequence 20, Appl	626	28	58.3	2843	2	US-08-450-582-7	Sequence 7, Appl1
554	28	58.3	998	2	US-08-368-776A-2	Sequence 2, Appl1	627	28	58.3	2843	2	US-08-449-731-2	Sequence 2, Appl1
555	28	58.3	998	2	US-09-949-016-6501	Sequence 6501, Ap	628	28	58.3	2843	2	US-10-092-138A-30	Sequence 30, Appl
556	28	58.3	998	4	PCT-US95-04681-17	Sequence 17, Appl	629	28	58.3	2843	2	US-09-538-092-1007	Sequence 1007, Ap
557	28	58.3	998	4	PCT-US95-04681-20	Sequence 20, Appl	630	28	58.3	2843	2	US-08-681-219A-30	Sequence 30, Appl
558	28	58.3	998	4	PCT-US96-00419-2	Sequence 2, Appl	631	28	58.3	2973	1	US-08-821-355A-7	Sequence 7, Appl1
559	28	58.3	1005	4	US-09-949-016-6968	Sequence 6968, Ap	632	28	58.3	2973	1	US-08-821-355A-7	Sequence 7, Appl1
560	28	58.3	1005	2	US-09-949-016-9901	Sequence 9901, Ap	633	28	58.3	2973	1	US-09-003-687A-7	Sequence 7, Appl1
561	28	58.3	1005	2	US-09-949-016-10620	Sequence 10620, A	634	28	58.3	2973	2	US-09-136-605-7	Sequence 7, Appl1
562	28	58.3	1011	1	US-08-162-809-12	Sequence 12, Appl	635	28	58.3	3177	1	US-08-477-451-4	Sequence 4, Appl1
563	28	58.3	1104	2	US-08-222-616-36	Sequence 36, Appl	636	28	58.3	3177	2	US-09-538-092-1136	Sequence 1136, Ap
564	28	58.3	1104	2	US-08-446-648-36	Sequence 36, Appl	637	28	58.3	3729	2	US-08-804-227C-4	Sequence 4, Appl1
565	28	58.3	1104	2	US-09-982-610-36	Sequence 36, Appl	638	28	57.3	823	1	US-08-248-766A-19339	Sequence 19339, A
566	28	58.3	1104	4	PCT-US95-04228-36	Sequence 36, Appl	639	27	56.2	15	2	US-08-553-287A-55	Sequence 55, Appl
567	28	58.3	1115	2	US-10-012-389A-58	Sequence 58, Appl	640	27	56.2	41	2	US-09-441-992-55	Sequence 55, Appl
568	28	58.3	1115	2	US-10-012-389A-58	Sequence 58, Appl	641	27	56.2	41	2	US-09-270-767-57140	Sequence 57140, A
569	28	58.3	1115	2	US-10-006-768A-58	Sequence 58, Appl	642	27	56.2	45	2	US-09-902-540-16522	Sequence 16522, A
570	28	58.3	1115	2	US-10-015-671A-58	Sequence 58, Appl	643	27	56.2	56	2	US-09-270-767-13867	Sequence 13867, A
571	28	58.3	1115	2	US-10-015-393A-58	Sequence 58, Appl	644	27	56.2	57	1	US-08-241-853-3	Sequence 3, Appl1
572	28	58.3	1115	2	US-10-011-833A-58	Sequence 58, Appl	645	27	56.2	57	1	US-08-917-1-3	Sequence 3, Appl1
573	28	58.3	1115	2	US-10-006-041A-58	Sequence 58, Appl	646	27	56.2	64	2	US-09-248-766A-22717	Sequence 22717, A
574	28	58.3	1115	2	US-10-012-064A-58	Sequence 58, Appl	647	27	56.2	65	2	US-09-134-000C-5463	Sequence 5463, Ap
575	28	58.3	1118	1	US-08-724-354D-2	Sequence 2, Appl1	648	27	56.2	66	2	US-09-270-767-11370	Sequence 11370, A
576	28	58.3	1118	2	US-09-270-984A-2	Sequence 2, Appl1	649	27	56.2	66	2	US-09-270-767-6586	Sequence 6586, A
577	28	58.3	1140	2	US-09-949-016-10116	Sequence 10116, A	650	27	56.2	93	2	US-09-270-767-61859	Sequence 61859, A
578	28	58.3	1182	2	US-09-041-886-21	Sequence 21, Appl	651	27	56.2	94	1	US-08-341-843B-30	Sequence 30, Appl
579	28	58.3	1194	2	US-09-092-508-2	Sequence 2, Appl1	652	27	56.2	94	1	US-08-427-497B-35	Sequence 35, Appl
580	28	58.3	1194	2	US-09-435-115-2	Sequence 2, Appl1	653	27	56.2	96	1	US-08-341-843B-6	Sequence 6, Appl1
581	28	58.3	1194	2	US-09-069-023-26	Sequence 26, Appl	654	27	56.2	96	1	US-08-427-497B-11	Sequence 11, Appl
582	28	58.3	1194	2	US-09-098-310-2	Sequence 2, Appl1	655	27	56.2	99	2	US-08-466-368-10	Sequence 10, Appl
583	28	58.3	1194	2	US-09-538-092-825	Sequence 825, App	656	27	56.2	99	2	US-08-470-998-7	Sequence 7, Appl1
584	28	58.3	1194	2	US-09-949-016-6030	Sequence 6030, Ap	657	27	56.2	99	2	US-09-621-976-4323	Sequence 4323, Ap
585	28	58.3	1196	2	US-09-949-016-10065	Sequence 10065, A	658	27	56.2	99	2	US-08-328-500-15	Sequence 15, Appl
586	28	58.3	1196	2	US-09-949-016-10066	Sequence 10066, A	659	27	56.2	102	2	US-09-605-703B-1358	Sequence 1358, Ap
587	28	58.3	1205	2	US-09-092-508-16	Sequence 16, Appl	660	27	56.2	103	2	US-09-513-999C-4915	Sequence 4915, Ap
588	28	58.3	1205	2	US-09-435-115-16	Sequence 16, Appl	661	27	56.2	104	2	US-08-99C-338-24	Sequence 24, Appl
589	28	58.3	1224	2	US-09-902-540-16312	Sequence 16312, A	662	27	56.2	104	2	US-09-556-972-24	Sequence 24, Appl
590	28	58.3	1237	2	US-09-949-016-6842	Sequence 6842, Ap	663	27	56.2	107	2	US-09-621-976-3893	Sequence 3893, Ap
591	28	58.3	1239	2	US-09-949-016-10063	Sequence 10063, A	664	27	56.2	113	2	US-09-377-285B-65	Sequence 65, Appl
592	28	58.3	1239	2	US-09-949-016-10064	Sequence 10064, A	665	27	56.2	113	2	US-10-192-381-65	Sequence 65, Appl
593	28	58.3	1276	1	US-08-222-616-24	Sequence 24, Appl	666	27	56.2	119	1	US-08-219-327B-3	Sequence 3, Appl1
594	28	58.3	1276	2	US-08-446-648-24	Sequence 24, Appl	667	27	56.2	119	2	US-08-477-347-14	Sequence 14, Appl
595	28	58.3	1276	2	US-09-982-610-24	Sequence 24, Appl	668	27	56.2	119	2	US-08-476-862-5	Sequence 5, Appl1
596	28	58.3	1276	4	PCT-US95-04228-34	Sequence 24, Appl	669	27	56.2	119	2	US-08-468-560C-3	Sequence 3, Appl1
597	28	58.3	1276	4	PCT-US95-04228-34	Sequence 24, Appl	670	27	56.2	119	2	US-08-828-663A-15	Sequence 15, Appl
598	28	58.3	1501	2	US-09-976-594-531	Sequence 531, App	671	27	56.2	119	2	US-09-800-909-5	Sequence 5, Appl1
599	28	58.3	1501	2	US-08-447-464-3	Sequence 3, Appl1	672	27	56.2	119	2	US-09-800-908-14	Sequence 14, Appl
600	28	58.3	1770	2	US-08-716-679-3	Sequence 3, Appl1	673	27	56.2	119	2	US-09-884-987-3	Sequence 3, Appl1
601	28	58.3	1911	1	US-10-144-198-44	Sequence 44, Appl	674	27	56.2	122	2	US-08-476-862-5	Sequence 3887, Ap
602	28	58.3	1911	1	US-08-348-006B-5	Sequence 4, Appl	675	27	56.2	122	2	US-10-104-047-3887	Sequence 3887, Ap
603	28	58.3	1911	1	US-08-800-825A-5	Sequence 5, Appl1	676	27	56.2	128	2	US-09-180-100-9	Sequence 9, Appl1
604	28	58.3	1911	1	US-09-158-657-5	Sequence 5, Appl1	677	27	56.2	128	2	US-09-949-016-713-9	Sequence 9, Appl1
605	28	58.3	1911	4	PCT-US94-1016-5	Sequence 5, Appl1	678	27	56.2	130	2	US-09-270-767-50662	Sequence 50662, A
606	28	58.3	2221	2	US-10-144-198-30	Sequence 30, Appl	679	27	56.2	134	2	US-09-270-767-34664	Sequence 34664, A
607	28	58.3	2265	2	US-09-269-446D-44	Sequence 44, Appl	680	27	56.2	134	2	US-09-270-767-34664	Sequence 34664, A
608	28	58.3	2308	1	US-08-149-097D-36	Sequence 36, Appl	681	27	56.2	139	2	US-09-857-076-13	Sequence 13, Appl
609	28	58.3	2308	1	US-08-015-973-1	Sequence 1, Appl1	682	27	56.2	141	2	US-09-205-658-13	Sequence 13, Appl
610	28	58.3	2308	1	US-08-448-164-1	Sequence 1, Appl1	683	27	56.2	141	2	US-09-583-110-2819	Sequence 2819, Ap
611	28	58.3	2308	2	US-08-081-929-2	Sequence 2, Appl1	684	27	56.2	141	2	US-09-107-433-4958	Sequence 4958, Ap

685	27	56.2	143	2	US-09-180-100-10	Sequence 10, Appl	758	27	56.2	309	2	US-09-556-972-22	Sequence 22, Appl
686	27	56.2	143	2	US-09-949-713-10	Sequence 10, Appl	759	27	56.2	313	2	US-09-413-814-9	Sequence 9, Appl
687	27	56.2	144	2	US-09-180-100-21	Sequence 21, Appl	760	27	56.2	314	1	US-08-444-231-19	Sequence 19, Appl
688	27	56.2	144	2	US-09-949-713-21	Sequence 21, Appl	761	27	56.2	314	1	US-08-152-433A-19	Sequence 19, Appl
689	27	56.2	148	2	US-09-270-767-13805	Sequence 33805, A	762	27	56.2	314	4	PCT-US95-17083-44	Sequence 4, Appl
690	27	56.2	148	2	US-09-270-767-49022	Sequence 49022, A	763	27	56.2	319	2	US-08-836-0785A-44	Sequence 230, Appl
691	27	56.2	148	2	US-09-270-767-57157	Sequence 57157, A	764	27	56.2	319	2	US-08-635-886C-230	Sequence 230, Appl
692	27	56.2	152	2	US-09-270-767-42700	Sequence 42700, A	765	27	56.2	319	2	US-08-974-690C-230	Sequence 230, Appl
693	27	56.2	157	2	US-09-180-100-15	Sequence 15, Appl	766	27	56.2	321	2	US-09-248-796A-14331	Sequence 14331, A
694	27	56.2	157	2	US-09-949-713-15	Sequence 15, Appl	767	27	56.2	322	2	US-09-198-452A-226	Sequence 226, App
695	27	56.2	159	2	US-09-180-100-23	Sequence 23, Appl	768	27	56.2	328	2	US-09-248-796A-15564	Sequence 15564, A
696	27	56.2	159	2	US-09-508-691-5	Sequence 5, Appl	769	27	56.2	331	2	US-09-086-483A-3	Sequence 3, Appl
697	27	56.2	159	2	US-09-949-713-23	Sequence 23, Appl	770	27	56.2	331	2	US-09-580-212-3	Sequence 3, Appl
698	27	56.2	162	2	US-09-270-767-49086	Sequence 49086, A	771	27	56.2	331	2	US-09-252-991A-29799	Sequence 29799, A
699	27	56.2	167	2	US-08-828-683A-22	Sequence 22, Appl	772	27	56.2	331	2	US-09-769-402-3	Sequence 3, Appl
700	27	56.2	173	2	US-09-357-251-8	Sequence 8, Appl	773	27	56.2	335	1	US-08-219-237B-2	Sequence 2, Appl
701	27	56.2	175	2	US-09-902-540-16214	Sequence 16214, A	774	27	56.2	335	1	US-08-409-338-1	Sequence 1, Appl
702	27	56.2	178	2	US-09-252-991A-20939	Sequence 20939, A	775	27	56.2	335	2	US-08-815-469-6	Sequence 6, Appl
703	27	56.2	178	2	US-09-267-963D-33	Sequence 33, Appl	776	27	56.2	335	2	US-09-290-640-2	Sequence 2, Appl
704	27	56.2	180	2	US-09-710-279-1588	Sequence 1588, Ap	777	27	56.2	335	2	US-09-006-353A-7	Sequence 7, Appl
705	27	56.2	180	2	US-09-893-737-168	Sequence 168, App	778	27	56.2	335	2	US-08-468-560C-2	Sequence 2, Appl
706	27	56.2	184	2	US-09-867-753-2	Sequence 2, Appl	779	27	56.2	335	2	US-09-180-100-20	Sequence 20, Appl
707	27	56.2	185	2	US-09-270-767-57514	Sequence 57514, A	780	27	56.2	335	2	US-09-565-918-3	Sequence 3, Appl
708	27	56.2	189	2	US-09-270-767-41893	Sequence 41893, A	781	27	56.2	335	2	US-09-573-986-7	Sequence 7, Appl
709	27	56.2	192	2	US-09-134-001C-4652	Sequence 4652, Ap	782	27	56.2	335	2	US-09-665-615B-2	Sequence 2, Appl
710	27	56.2	193	2	US-09-489-039A-10031	Sequence 10031, A	783	27	56.2	335	2	US-09-557-908-6	Sequence 6, Appl
711	27	56.2	193	2	US-09-270-767-39645	Sequence 39645, A	784	27	56.2	335	2	US-09-874-138-4	Sequence 4, Appl
712	27	56.2	193	2	US-09-270-767-54862	Sequence 54862, A	785	27	56.2	335	2	US-09-333-966-6	Sequence 6, Appl
713	27	56.2	204	2	US-09-248-796A-23484	Sequence 23484, A	786	27	56.2	335	2	US-09-949-016-5877	Sequence 5877, Ap
714	27	56.2	207	2	US-08-996-338-23	Sequence 23, Appl	787	27	56.2	335	2	US-09-565-009B-4	Sequence 4, Appl
715	27	56.2	207	2	US-09-556-972-33	Sequence 33, Appl	788	27	56.2	335	2	US-10-175-902-3	Sequence 3, Appl
716	27	56.2	209	2	US-09-761-534A-12	Sequence 12, Appl	789	27	56.2	335	2	US-09-995-938A-8	Sequence 8, Appl
717	27	56.2	212	2	US-09-270-767-33190	Sequence 33190, A	790	27	56.2	335	2	US-09-884-887-2	Sequence 10, Appl
718	27	56.2	212	2	US-09-902-540-15176	Sequence 15176, A	791	27	56.2	335	2	US-09-314-889-6	Sequence 6, Appl
719	27	56.2	219	2	US-08-974-022-45	Sequence 45, Appl	792	27	56.2	335	2	US-09-949-713-20	Sequence 20, Appl
720	27	56.2	219	2	US-08-795-445A-45	Sequence 45, Appl	793	27	56.2	335	4	PCT-US95-17083-2	Sequence 2, Appl
721	27	56.2	219	2	US-08-795-447A-45	Sequence 45, Appl	794	27	56.2	335	4	US-09-198-452A-662	Sequence 662, App
722	27	56.2	219	2	US-08-974-186-45	Sequence 45, Appl	795	27	56.2	336	2	US-09-438-185A-625	Sequence 625, App
723	27	56.2	219	2	US-08-795-945B-45	Sequence 45, Appl	796	27	56.2	336	2	US-09-438-185A-212	Sequence 212, App
724	27	56.2	219	2	US-08-706-945D-131	Sequence 131, App	797	27	56.2	342	2	US-08-978-456-2	Sequence 2, Appl
725	27	56.2	219	2	US-08-577-788C-45	Sequence 45, Appl	798	27	56.2	342	2	US-09-369-700-2	Sequence 2, Appl
726	27	56.2	225	2	US-10-104-047-3107	Sequence 3107, Ap	799	27	56.2	344	2	US-09-489-039A-285	Sequence 285, App
727	27	56.2	245	2	US-09-248-796A-19933	Sequence 19933, A	800	27	56.2	344	2	US-09-489-039A-11854	Sequence 11854, A
728	27	56.2	255	2	US-09-543-681A-6904	Sequence 6904, Ap	801	27	56.2	350	2	US-09-902-540-10961	Sequence 10961, A
729	27	56.2	257	2	US-09-489-039A-7196	Sequence 7196, Ap	802	27	56.2	352	2	US-09-270-767-42603	Sequence 42603, A
730	27	56.2	260	2	US-09-489-039A-11113	Sequence 11113, A	803	27	56.2	357	2	US-09-270-767-42603	Sequence 6082, Ap
731	27	56.2	266	2	US-09-248-796A-19766	Sequence 19766, A	804	27	56.2	359	2	US-09-543-681A-6082	Sequence 61909, A
732	27	56.2	270	2	US-09-489-039A-12261	Sequence 12261, A	805	27	56.2	359	2	US-09-270-767-41909	Sequence 41909, A
733	27	56.2	276	2	US-09-270-767-44260	Sequence 44260, A	806	27	56.2	360	2	US-09-180-100-11	Sequence 11, Appl
734	27	56.2	276	2	US-09-248-796A-24171	Sequence 24171, A	807	27	56.2	360	2	US-09-538-092-547	Sequence 547, App
735	27	56.2	276	2	US-09-949-016-9958	Sequence 9958, Ap	808	27	56.2	360	2	US-09-949-713-11	Sequence 11, Appl
736	27	56.2	281	2	US-09-527-236A-3	Sequence 3, Appl	809	27	56.2	369	2	US-09-538-092-1166	Sequence 1166, Ap
737	27	56.2	281	2	US-09-756-854-3	Sequence 3, Appl	810	27	56.2	374	2	US-09-252-991A-29399	Sequence 29399, A
738	27	56.2	281	2	US-10-041-574-3	Sequence 3, Appl	811	27	56.2	376	2	US-09-180-100-22	Sequence 22, Appl
739	27	56.2	281	2	US-09-095-094-3	Sequence 3, Appl	812	27	56.2	376	2	US-09-949-713-22	Sequence 11, Appl
740	27	56.2	283	1	US-08-723-886-3	Sequence 3, Appl	813	27	56.2	378	2	US-08-158-735A-11	Sequence 22, Appl
741	27	56.2	283	1	US-08-972-902-8	Sequence 8, Appl	814	27	56.2	378	2	US-09-716-865-22	Sequence 22, Appl
742	27	56.2	283	2	US-09-466-257A-8	Sequence 8, Appl	815	27	56.2	378	2	US-09-270-767-46290	Sequence 46290, A
743	27	56.2	283	2	US-09-466-257A-10	Sequence 10, Appl	816	27	56.2	378	2	US-09-107-532A-3904	Sequence 3904, Ap
744	27	56.2	283	2	US-08-424-797A-3	Sequence 3, Appl	817	27	56.2	380	2	US-09-248-796A-19948	Sequence 19948, A
745	27	56.2	283	2	US-09-520-207-8	Sequence 8, Appl	818	27	56.2	387	2	US-09-134-001C-5036	Sequence 5036, Ap
746	27	56.2	283	2	US-10-136-253-8	Sequence 8, Appl	819	27	56.2	387	2	US-09-949-016-7720	Sequence 7720, Ap
747	27	56.2	286	2	US-09-583-110-2939	Sequence 2939, Ap	820	27	56.2	390	2	US-09-902-540-12958	Sequence 12958, A
748	27	56.2	287	2	US-08-311-731A-2332	Sequence 2332, App	821	27	56.2	392	1	US-08-886-152-1	Sequence 1, Appl
749	27	56.2	287	2	US-09-107-433-5005	Sequence 5005, Ap	822	27	56.2	392	2	US-09-156-422-1	Sequence 1, Appl
750	27	56.2	294	2	US-09-270-767-59588	Sequence 59588, A	823	27	56.2	397	2	US-09-489-039A-8962	Sequence 8962, Ap
751	27	56.2	297	2	US-09-489-039A-7396	Sequence 7396, Ap	824	27	56.2	403	2	US-09-252-991A-17989	Sequence 17989, A
752	27	56.2	298	2	US-09-489-039A-12886	Sequence 12886, A	825	27	56.2	403	2	US-09-270-767-42237	Sequence 42237, A
753	27	56.2	301	2	US-09-252-991A-18879	Sequence 18879, A	826	27	56.2	407	2	US-09-198-452A-805	Sequence 805, App
754	27	56.2	306	2	US-09-252-991A-20009	Sequence 20009, Appl	827	27	56.2	407	2	US-09-438-185A-757	Sequence 757, App
755	27	56.2	308	2	US-09-369-247-60	Sequence 60, Appl	828	27	56.2	416	2	US-09-976-594-810	Sequence 810, App
756	27	56.2	308	2	US-10-062-548-60	Sequence 60, Appl	829	27	56.2	416	2	US-09-949-016-6385	Sequence 6385, Ap
757	27	56.2	309	2	US-08-996-338-22	Sequence 22, Appl	830	27	56.2	417	2	US-09-433-241A-10	Sequence 10, Appl

831	27	56.2	429	2	US-09-489-039A-13930	Sequence 13930, A	904	27	56.2	669	2	US-09-013-895A-3	Sequence 3, Appl1
832	27	56.2	434	2	US-09-433-241A-14	Sequence 14, Appl1	905	27	56.2	669	2	US-09-448-868-3	Sequence 3, Appl1
833	27	56.2	435	2	US-09-433-241A-12	Sequence 12, Appl1	906	27	56.2	669	2	US-10-226-296-3	Sequence 3, Appl1
834	27	56.2	436	2	US-09-721-870-111	Sequence 11, App	907	27	56.2	670	2	US-09-248-796A-17669	Sequence 17669, A
835	27	56.2	437	2	US-09-543-681A-7455	Sequence 7455, Ap	908	27	56.2	684	2	US-09-823-240A-9	Sequence 9, Appl1
836	27	56.2	443	2	US-09-949-016-11613	Sequence 11613, A	909	27	56.2	708	2	US-09-949-016-7574	Sequence 7574, Ap
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838	27	56.2	448	2	US-09-438-185A-1016	Sequence 1016, Ap	911	27	56.2	710	2	US-09-270-767-43621	Sequence 43621, A
839	27	56.2	450	2	US-09-949-016-8611	Sequence 8611, Ap	912	27	56.2	718	1	US-08-190-802A-44	Sequence 44, Appl1
840	27	56.2	450	2	US-09-949-016-9597	Sequence 9597, Ap	913	27	56.2	718	2	US-08-477-346-44	Sequence 44, Appl1
841	27	56.2	468	2	US-09-292-097-2	Sequence 2, Appl1	914	27	56.2	718	2	US-08-487-072A-44	Sequence 44, Appl1
842	27	56.2	468	2	US-09-933-561-2	Sequence 2, Appl1	915	27	56.2	718	2	US-08-487-072A-44	Sequence 44, Appl1
843	27	56.2	495	2	US-09-949-016-9023	Sequence 9023, Ap	916	27	56.2	731	2	US-09-252-991A-25890	Sequence 25890, A
844	27	56.2	502	2	US-09-904-615-69	Sequence 69, Appl1	917	27	56.2	731	2	US-09-232-338-10	Sequence 10, Appl1
845	27	56.2	502	2	US-10-054-988-69	Sequence 69, Appl1	918	27	56.2	741	2	US-09-543-681A-8128	Sequence 8128, Ap
846	27	56.2	505	2	US-09-270-767-11697	Sequence 41697, A	919	27	56.2	751	2	US-09-540-236-2921	Sequence 2921, Ap
847	27	56.2	508	2	US-09-811-469-7	Sequence 7, Appl1	920	27	56.2	764	2	US-09-902-140-14495	Sequence 6, Appl1
848	27	56.2	508	2	US-10-370-659-7	Sequence 7, Appl1	921	27	56.2	766	4	PCT-US94-00198-6	Sequence 6, Appl1
849	27	56.2	512	2	US-09-248-796A-15186	Sequence 15186, A	922	27	56.2	771	1	US-07-923-976-6	Sequence 18066, A
850	27	56.2	514	2	US-09-710-279-2682	Sequence 2682, Ap	923	27	56.2	778	2	US-09-248-796A-18066	Patent No. 5422248
851	27	56.2	518	2	US-09-134-001C-4451	Sequence 4451, Ap	924	27	56.2	783	6	5422248-2	Sequence 15074, A
852	27	56.2	520	2	US-09-527-073-2	Sequence 2, Appl1	925	27	56.2	835	1	US-09-248-796A-15074	Sequence 4, Appl1
853	27	56.2	521	2	US-08-996-338-20	Sequence 20, Appl1	926	27	56.2	836	1	US-07-923-976-4	Sequence 62, Appl1
854	27	56.2	521	2	US-09-556-972-20	Sequence 20, Appl1	927	27	56.2	854	2	US-09-487-558B-62	Sequence 16013, A
855	27	56.2	527	2	US-09-489-039A-9695	Sequence 9695, Ap	928	27	56.2	862	2	US-09-248-796A-16013	Sequence 8, Appl1
856	27	56.2	538	2	US-09-252-991A-24079	Sequence 24079, A	929	27	56.2	863	1	US-07-923-976-8	Sequence 20365, A
857	27	56.2	541	1	US-08-604-333-2	Sequence 2, Appl1	930	27	56.2	873	2	US-09-248-796A-20165	Sequence 13902, A
858	27	56.2	541	2	US-09-110-618-2	Sequence 28, Appl1	931	27	56.2	884	2	US-09-902-540-13902	Sequence 2474, Ap
859	27	56.2	541	2	US-09-173-151A-28	Sequence 2, Appl1	932	27	56.2	897	2	US-10-104-047-2474	Sequence 2, Appl1
860	27	56.2	541	2	US-09-578-178-2	Sequence 2, Appl1	933	27	56.2	928	2	US-09-336-946B-2	Sequence 4, Appl1
861	27	56.2	541	2	US-09-577-806-2	Sequence 2, Appl1	934	27	56.2	928	2	US-09-336-946B-4	Sequence 58, Appl1
862	27	56.2	541	2	US-09-621-502-4	Sequence 4, Appl1	935	27	56.2	928	2	US-09-993-170-58	Sequence 30698, A
863	27	56.2	541	2	US-09-949-002-360	Sequence 360, App	936	27	56.2	985	2	US-09-252-991A-30698	Sequence 8664, Ap
864	27	56.2	545	2	US-09-248-796A-18865	Sequence 18865, A	937	27	56.2	1022	2	US-09-949-016-8864	Sequence 9041, Ap
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866	27	56.2	551	2	US-08-688-988-3736	Sequence 37, Appl1	939	27	56.2	1043	1	US-08-562-737-46	Sequence 4, Appl1
867	27	56.2	551	2	US-09-540-236-2436	Sequence 2436, Ap	940	27	56.2	1043	2	US-08-724-354D-4	Sequence 15476, A
868	27	56.2	565	1	US-08-357-533A-9	Sequence 9, Appl1	941	27	56.2	1060	2	US-09-270-798A-15476	Sequence 95, Appl1
869	27	56.2	565	1	US-08-459-001-9	Sequence 9, Appl1	942	27	56.2	1120	2	US-09-792-024-95	Sequence 18, Appl1
870	27	56.2	565	2	US-08-459-951-9	Sequence 9, Appl1	943	27	56.2	1132	1	US-08-446-038B-18	Sequence 18, Appl1
871	27	56.2	567	1	US-08-361-873A-2	Sequence 2, Appl1	944	27	56.2	1132	1	US-08-446-038B-18	Sequence 18, Appl1
872	27	56.2	567	1	US-08-483-926A-1	Sequence 1, Appl1	945	27	56.2	1132	1	US-08-446-010B-18	Sequence 18, Appl1
873	27	56.2	567	1	US-08-854-768-1	Sequence 1, Appl1	946	27	56.2	1132	1	US-08-805-445-18	Sequence 18, Appl1
874	27	56.2	567	1	US-08-445-520B-9	Sequence 9, Appl1	947	27	56.2	1132	1	US-08-064-067D-18	Sequence 18, Appl1
875	27	56.2	567	1	US-08-737-045-1	Sequence 1, Appl1	948	27	56.2	1132	1	US-09-066-208-18	Sequence 18, Appl1
876	27	56.2	567	2	US-08-451-946B-8	Sequence 8, Appl1	949	27	56.2	1135	2	US-08-574-959A-7	Sequence 7, Appl1
877	27	56.2	567	2	US-08-446-938B-8	Sequence 8, Appl1	950	27	56.2	1142	2	US-08-097-997A-11	Sequence 11, Appl1
878	27	56.2	567	2	US-08-311-703A-8	Sequence 8, Appl1	951	27	56.2	1142	2	US-08-665-574C-11	Sequence 11, Appl1
879	27	56.2	567	2	US-08-446-939B-8	Sequence 8, Appl1	952	27	56.2	1142	2	US-08-946-994A-11	Sequence 21, Appl1
880	27	56.2	567	2	US-09-183-543-8	Sequence 8, Appl1	953	27	56.2	1142	2	US-08-771-161A-211	Sequence 14, Appl1
881	27	56.2	567	2	US-08-446-936A-8	Sequence 8, Appl1	954	27	56.2	1153	1	US-08-097-997A-14	Sequence 14, Appl1
882	27	56.2	567	2	US-08-446-936A-8	Sequence 8, Appl1	955	27	56.2	1153	2	US-08-665-574C-14	Sequence 14, Appl1
883	27	56.2	567	2	US-09-239-864A-11	Sequence 11, Appl1	956	27	56.2	1153	2	US-08-946-994A-14	Sequence 14, Appl1
884	27	56.2	567	2	US-09-878-905-11	Sequence 11, Appl1	957	27	56.2	1153	2	US-08-946-994A-14	Sequence 14, Appl1
885	27	56.2	567	2	US-09-267-963D-36	Sequence 36, Appl1	958	27	56.2	1154	1	US-08-357-598-7	Sequence 7, Appl1
886	27	56.2	567	4	PCT-US92-09326-4	Sequence 4, Appl1	959	27	56.2	1154	1	US-08-446-010B-24	Sequence 24, Appl1
887	27	56.2	572	2	US-09-771-161A-120	Sequence 120, App	960	27	56.2	1154	4	US-09-003-289-7	Sequence 7, Appl1
888	27	56.2	578	2	US-09-489-039A-10599	Sequence 10599, A	961	27	56.2	1154	4	PCT-US95-16435-7	Sequence 24669, A
889	27	56.2	578	2	US-09-248-796A-14802	Sequence 14802, A	962	27	56.2	1218	2	US-09-252-991A-24869	Sequence 7874, Ap
890	27	56.2	590	1	US-08-448-196A-9	Sequence 9, Appl1	963	27	56.2	1233	2	US-09-328-352-7874	Sequence 2, Appl1
891	27	56.2	595	2	US-09-489-039A-13433	Sequence 13433, A	964	27	56.2	1233	1	US-08-241-853-2	Sequence 2, Appl1
892	27	56.2	602	1	US-08-419-652-6	Sequence 6, Appl1	965	27	56.2	1237	1	US-08-850-917-2	Sequence 21, Appl1
893	27	56.2	602	1	US-09-569-037-6	Sequence 6, Appl1	966	27	56.2	1260	2	US-08-506-236B-21	Sequence 64, Appl1
894	27	56.2	612	2	US-09-248-796A-20967	Sequence 20967, A	967	27	56.2	1321	1	US-08-317-310A-64	Sequence 14121, A
895	27	56.2	622	2	US-10-132-556A-2	Sequence 2, Appl1	968	27	56.2	1503	2	US-08-976-255-14	Sequence 43621, A
896	27	56.2	640	1	US-08-671-978A-10	Sequence 10, Appl1	969	27	56.2	1896	2	US-09-949-016-9508	Sequence 44, Appl1
897	27	56.2	641	2	US-09-919-039-146	Sequence 146, App	970	27	56.2	3200	1	US-08-477-451-8	Sequence 44, Appl1
898	27	56.2	642	2	US-09-761-534A-10	Sequence 10, Appl1	971	27	56.2	4019	2	US-09-854-133-425	Sequence 425, App
899	27	56.2	659	2	US-09-562-737-19	Sequence 19, Appl1	972	27	56.2	4126	2	US-09-953-036-4	Sequence 4, Appl1
900	27	56.2	668	2	US-09-811-469-6	Sequence 6, Appl1	973	27	56.2	4968	2	US-09-424-783-5	Sequence 5, Appl1
901	27	56.2	669	2	US-10-370-659-6	Sequence 6, Appl1	974	27	56.2	5518	2	US-09-953-036-2	Sequence 2, Appl1
902	27	56.2	669	1	US-08-357-533A-8	Sequence 8, Appl1	975	27	55.2	171	2	US-09-270-767-44163	Sequence 34163, A
903	27	56.2	669	2	US-08-459-009-8	Sequence 8, Appl1	976	26.5	55.2	171	2	US-09-270-767-49380	Sequence 49380, A

977 26.5 55.2 905 2 US-09-360-186-3 Sequence 3, Appl1
978 26.5 55.2 905 2 US-09-864-680A-3 Sequence 3, Appl1
979 26 54.2 9 1 US-08-787-547-102 Sequence 102, App
980 26 54.2 9 2 US-08-159-339A-248 Sequence 248, App
981 26 54.2 34 2 US-09-324-455-19 Sequence 19, Appl
982 26 54.2 38 2 US-09-205-258-982 Sequence 982, App
983 26 54.2 38 2 US-10-004-860-982 Sequence 982, App
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985 26 54.2 64 2 US-09-248-796A-26931 Sequence 26931, A
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988 26 54.2 65 2 US-09-107-433-3119 Sequence 3119, A
989 26 54.2 66 2 US-09-621-976-5979 Sequence 5979, Ap
990 26 54.2 67 2 US-09-248-796A-22839 Sequence 22839, A
991 26 54.2 73 2 US-09-248-796A-22839 Sequence 22839, A
992 26 54.2 78 2 US-09-270-767-36289 Sequence 36289, A
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995 26 54.2 84 2 US-09-247-155-119 Sequence 119, App
996 26 54.2 84 2 US-09-621-976-7012 Sequence 7012, App
997 26 54.2 84 2 US-09-270-767-43718 Sequence 43718, A
998 26 54.2 84 2 US-09-903-190-119 Sequence 119, App
999 26 54.2 88 2 US-09-583-110-4548 Sequence 4548, Ap
1000 26 54.2 88 2 US-09-583-110-4548 Sequence 4548, Ap

ALIGNMENTS

RESULT 1
US-08-159-339A-73
Sequence 73, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esteban
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 73:

SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-73

Query Match 100.0%; Score 48; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 1 INDIIIECV 9
Db 1 INDIIIECV 9

RESULT 2
US-08-197-484-72
Sequence 72, Application US/08197484
Patent No. 6419931
GENERAL INFORMATION:
APPLICANT: VITIELLO, Maria A.
APPLICANT: CHESTNUT, Robert W.
APPLICANT: SETTE, Alessandro D.
APPLICANT: CELIS, Esteban
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
TITLE OF INVENTION: CTL IMMUNITY
NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Knourie and Crew
STREET: Steuart Street Tower, One Market Plaza
CITY: San Francisco
STATE: California
COUNTRY: US
ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/197,484
FILING DATE: 16-FEB-1994
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935,811
FILING DATE: 26-AUG-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/827,682
FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W.
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 623-6793
TELEFAX: (206) 623-6793
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
US-08-197-484-72

Query Match 100.0%; Score 48; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 2 IHDIILLCV 10

RESULT 3

US-09-601-729-273
Sequence 273, Application US/09601729
Patent No. 6683052
GENERAL INFORMATION:
APPLICANT: THIAM, KADER
APPLICANT: AURIAULT, CLAUDE
APPLICANT: GRAS-MASSE, HELENE
APPLICANT: LOING, ESTELLE
APPLICANT: VERMAERDE, CLAUDIE
APPLICANT: GUILLET, JEAN GERARD
TITLE OF INVENTION: LIPOPEPTIDES CONTAINING AN INTERFERON FRAGMENT AND USES
FILE REFERENCE: USB-97-AU-IN
CURRENT APPLICATION NUMBER: US/09/601, 729
CURRENT FILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: PCT/FR99/00259
PRIOR FILING DATE: 1999-02-05
PRIOR APPLICATION NUMBER: 98 01439
PRIOR FILING DATE: 1998-02-06
NUMBER OF SEQ ID NOS: 281
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 273
LENGTH: 10
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-601-729-273

Query Match 100.0%; Score 48; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILLCV 9
DB 2 IHDIILLCV 10

RESULT 4

PCT-US95-02121-72
Sequence 72, Application PC/TUS9502121
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
TITLE OF INVENTION: CTL IMMUNITY
NUMBER OF SEQUENCES: 153
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/02121
FILING DATE: 16-FEB-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/197,484
FILING DATE: 16-FEB-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/935,811
FILING DATE: 26-AUG-1992

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
PRIOR APPLICATION DATA: US 07/827,682
FILING DATE: 29-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W.
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4PC
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 467-9600
TELEFAX: (415) 543-5043
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
PCT-US95-02121-72

Query Match 100.0%; Score 48; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.011;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILLCV 9
DB 2 IHDIILLCV 10

RESULT 5

US-09-980-523A-4
Sequence 4, Application US/09980523A
Patent No. 6783763
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN GERARD
APPLICANT: CONNAN, FRANCES
APPLICANT: FERRIS, ESTELLE
TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
FILE REFERENCE: WO1 AO INS
CURRENT APPLICATION NUMBER: US/09/980,523A
CURRENT FILING DATE: 2002-04-29
PRIOR APPLICATION NUMBER: PCT/FR00/01513
PRIOR FILING DATE: 2000-05-31
PRIOR APPLICATION NUMBER: FR 99/07012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 4
LENGTH: 30
TYPE: PRT
ORGANISM: Human Papillomavirus
US-09-980-523A-4

Query Match 100.0%; Score 48; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.033;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILLCV 9
DB 16 IHDIILLCV 24

RESULT 6

US-09-390-027-6

```
; Sequence 6, Application US/09390027
; Patent No. 6235523
; GENERAL INFORMATION:
; APPLICANT: GAJEWCZYK, Diane M.
; APPLICANT: PERSSON, Roy
; APPLICANT: YAO, Pei-Long
; APPLICANT: CAO, Shi-Xian
; APPLICANT: KLEIN, Michel H.
; APPLICANT: MARTAGLIA, James
; APPLICANT: MOINGEON, Philippe
; APPLICANT: ROVINSKI, Benjamin
; TITLE OF INVENTION: TREATMENT OF CERVICAL CANCER
; FILE REFERENCE: 1038-982 MIS-jb
; CURRENT APPLICATION NUMBER: US/09/390,027
; CURRENT FILING DATE: 1999-09-03
; EARLIER APPLICATION NUMBER: 60/099,291
; EARLIER FILING DATE: 1998-09-04
; NUMBER OF SEQ ID NOS: 12
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 6
; LENGTH: 59
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURES:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-09-390-027-6

Query Match      100.0%; Score 48; DB 2; Length 59;
Best Local Similarity 100.0%; Pred. No. 0.066;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 IHDIILRCV 9
Db      51 IHDIILRCV 59

RESULT 7
US-09-701-080C-18
; Sequence 18, Application US/09701080C
; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; CURRENT FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match      100.0%; Score 48; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.17;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 IHDIILRCV 9
Db      23 IHDIILRCV 31

RESULT 8
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
```

```
; APPLICANT: CHOPIN, JEANNINE
; APPLICANT: BOURGULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: WO1 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match      100.0%; Score 48; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.18;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      1 IHDIILRCV 9
Db      30 IHDIILRCV 38

RESULT 9
US-08-316-239B-3
; Sequence 3, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; TITLE OF INVENTION: Methods and a Diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 803-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
```

MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-3

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.18;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
Db 30 IHDIILECV 38

RESULT 10
US-08-316-239B-4
Sequence 4, Application US/08316239B
Patent No. 5679509
GENERAL INFORMATION:
APPLICANT: Wheeler, Cosette M.
APPLICANT: Patmeyer, Cheryl A.
TITLE OF INVENTION: Methods and a Diagnostic Aid for
TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
TITLE OF INVENTION: Cervical Cancer
NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Jagtland & Associates
STREET: 6126 Rocky Way Court
CITY: Centreville
STATE: VA
COUNTRY: USA
ZIP: 20120-3400
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/316,239B
FILING DATE: 30-SEP-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Jagtland, Ajay A.
REGISTRATION NUMBER: 35,205
REFERENCE/DOCKET NUMBER: UNME-0001
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703) 817-9453
TELEFAX: (703) 803-9387
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 162 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHETICAL: NO
US-08-316-239B-4

Query Match 100.0%; Score 48; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.18; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
Db 30 IHDIILECV 38

RESULT 11
US-08-860-165-14
Sequence 14, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John

APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCES: 17227/130
CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-14

Query Match 100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.19; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
Db 99 IHDIILECV 107

RESULT 12
US-09-359-382-14
Sequence 14, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FRAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCES: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU PNO157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 14
LENGTH: 172
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-14

Query Match 100.0%; Score 48; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.19; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
Db 99 IHDIILECV 107

RESULT 13
US-08-117-083-10
Sequence 10, Application US/08117083
Patent No. 5719054
GENERAL INFORMATION:
APPLICANT: Boursnell, Michael E.
APPLICANT: Inglis, Stephen C.

```

; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Walter H. Dregger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dregger, Walter H.
; REGISTRATION NUMBER: 24,150
; REFERENCE/DOCKET NUMBER: A-58783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 182 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
; FEATURE:
; NAME/KEY: Protein
; LOCATION: 1..182
; OTHER INFORMATION: /note="Xaa refers to stop codon in
; OTHER INFORMATION: the open reading frame."
; US-08-117-083-10

Query Match          100.0%; Score 48; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILCEV 9
DB 31 IHDIILCEV 39

RESULT 14
US-09-462-993-1
; Sequence 1, Application US/09462993
; Patent No. 6884786
; GENERAL INFORMATION:
; APPLICANT: KIENT, Marie-Paule
; APPLICANT: BALLOUL, Jean-Marc
; APPLICANT: BIZOUARNE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 01753-122
; CURRENT APPLICATION NUMBER: US/09/462,993
; CURRENT FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: Patentin Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
```

```

; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence:Derived from
; OTHER INFORMATION: human Papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone E6*TMF.
; US-09-462-993-1

Query Match          100.0%; Score 48; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.27;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILCEV 9
DB 58 IHDIILCEV 66

RESULT 15
US-08-860-165-10
; Sequence 10, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 17227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
; US-08-860-165-10

Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILCEV 9
DB 30 IHDIILCEV 38

RESULT 16
US-09-359-382-10
; Sequence 10, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
```



```
; SEQ ID NO 10
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-359-382-10
```

```
Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 IHDIILECV 9
Db      30 IHDIILECV 38
```

```
RESULT 17
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Patent No. 6428807
; GENERAL INFORMATION:
; APPLICANT: MACPARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 01727/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; PRIOR FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1
```

```
Query Match          100.0%; Score 48; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.3;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 IHDIILECV 9
Db      30 IHDIILECV 38
```

```
RESULT 18
US-09-485-885-4
; Sequence 4, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-4
```

```
Query Match          100.0%; Score 48; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.31;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 IHDIILECV 9
Db      136 IHDIILECV 144
```

```
RESULT 19
US-09-485-885-10
; Sequence 10, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-10
```

```
Query Match          100.0%; Score 48; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.33;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 IHDIILECV 9
Db      155 IHDIILECV 163
```

```
RESULT 20
US-09-485-885-6
; Sequence 6, Application US/09485885
; Patent No. 6342224
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/09/485,885
; PRIOR FILING DATE: 2000-02-18
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-09-485-885-6
```

```
Query Match          100.0%; Score 48; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.42;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

OY 1 IHDIILECV 9
 Db 136 IHDIILECV 144

RESULT 21
 US-09-485-885-14
 ; Sequence 14, Application US/09485885
 ; Patent No. 6342224
 ; GENERAL INFORMATION:
 ; APPLICANT: Bruck, Claudine
 ; APPLICANT: Cabezon Silva, Teresa
 ; APPLICANT: Delisse, Anne-Marie Eva Fernande
 ; APPLICANT: Gerard, Catherine Marie Ghislaine
 ; APPLICANT: Lombardo-Bencheikh, Angela
 ; TITLE OF INVENTION: Vaccine
 ; FILE REFERENCE: B45107
 ; CURRENT APPLICATION NUMBER: US/09/485,885
 ; CURRENT FILING DATE: 2000-02-18
 ; PRIOR APPLICATION NUMBER: PCT/EP98/05285
 ; PRIOR FILING DATE: 1998-08-17
 ; PRIOR APPLICATION NUMBER: GB 9717953.5
 ; PRIOR FILING DATE: 1997-08-22
 ; NUMBER OF SEQ ID NOS: 23
 ; SOFTWARE: PatsEQ for Windows Version 3.0
 ; SEQ ID NO 14
 ; LENGTH: 390
 ; TYPE: PRT
 ; ORGANISM: Homo sapien
 US-09-485-885-14

Query Match 100.0%; Score 48; DB 2; Length 390;
 Best Local Similarity 100.0%; Pred. No. 0.44;
 Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IHDIILECV 9
 Db 155 IHDIILECV 163

RESULT 22
 US-07-909-122-3
 ; Sequence 3, Application US/07909122
 ; Patent No. 5415995
 ; GENERAL INFORMATION:
 ; APPLICANT: SCHOOLNIK, GARY K.
 ; APPLICANT: PALESKY, JOEL M.
 ; TITLE OF INVENTION: DIAGNOSTIC PEPTIDES OF HUMAN PAPILLOMA
 ; TITLE OF INVENTION: VIRUS
 ; NUMBER OF SEQUENCES: 17
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: MORRISON & FOERSTER
 ; STREET: 755 Page Mill Road
 ; CITY: Palo Alto
 ; STATE: California
 ; COUNTRY: USA
 ; ZIP: 94304-1018
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: Patentin Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/07/909,122
 ; FILING DATE: 19920706
 ; CLASSIFICATION: 530
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: BENZ, WILLIAM H.
 ; REGISTRATION NUMBER: 25,952
 ; REFERENCE/DOCKET NUMBER: 28600-20105.01
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: (415) 813-5600

TELEFAX: (415) 494-0792
 ; TELEX: 706141
 ; INFORMATION FOR SEQ ID NO: 3:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 14 amino acids
 ; TYPE: AMINO ACID
 ; STRANDEDNESS: single
 ; TOPOLOGY: linear
 US-07-909-122-3

Query Match 91.7%; Score 44; DB 1; Length 14;
 Best Local Similarity 100.0%; Pred. No. 0.084;
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IHDIILEC 8
 Db 7 IHDIILEC 14

RESULT 23
 US-08-363-586-4
 ; Sequence 4, Application US/08363586
 ; Patent No. 5629161
 ; GENERAL INFORMATION:
 ; APPLICANT: Mueller, Martin
 ; APPLICANT: Giesmann, Lutz
 ; TITLE OF INVENTION: Use of HPV-16 E6 and E7-gene derived
 ; TITLE OF INVENTION: Peptides for the Diagnostic Purpose
 ; NUMBER OF SEQUENCES: 4
 ; CORRESPONDENCE ADDRESS:
 ; ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
 ; STREET: 1300 I Street, N.W.
 ; CITY: Washington
 ; STATE: D.C.
 ; COUNTRY: USA
 ; ZIP: 20005-3315
 ; COMPUTER READABLE FORM:
 ; MEDIUM TYPE: Floppy disk
 ; COMPUTER: IBM PC compatible
 ; OPERATING SYSTEM: PC-DOS/MS-DOS
 ; SOFTWARE: Patentin Release #1.0, Version #1.25
 ; CURRENT APPLICATION DATA:
 ; APPLICATION NUMBER: US/08/363,586
 ; FILING DATE: 23-DEC-1994
 ; CLASSIFICATION: 435
 ; PRIOR APPLICATION DATA:
 ; APPLICATION NUMBER: US 07/909,296
 ; FILING DATE: 09-JUL-1992
 ; APPLICATION NUMBER: EP 91111720.8
 ; FILING DATE: 13-JUL-1991
 ; ATTORNEY/AGENT INFORMATION:
 ; NAME: Wadler, Linda A.
 ; REGISTRATION NUMBER: 33,218
 ; REFERENCE/DOCKET NUMBER: 02481-1195-00000
 ; TELECOMMUNICATION INFORMATION:
 ; TELEPHONE: 202-408-4000
 ; TELEFAX: 202-408-4400
 ; INFORMATION FOR SEQ ID NO: 4:
 ; SEQUENCE CHARACTERISTICS:
 ; LENGTH: 30 amino acids
 ; TYPE: amino acid
 ; TOPOLOGY: linear
 ; MOLECULE TYPE: peptide
 US-08-363-586-4

Query Match 91.7%; Score 44; DB 1; Length 30;
 Best Local Similarity 100.0%; Pred. No. 0.18;
 Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

OY 1 IHDIILEC 8
 Db 23 IHDIILEC 30

```

RESULT 24
US-09-248-796A-15013
Sequence 15013, Application US/09248796A
Patent No. 6747137
GENERAL INFORMATION:
APPLICANT: Keith Weinstein et al
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: 107196.132
CURRENT APPLICATION NUMBER: US/09/248,796A
CURRENT FILING DATE: 1999-02-12
PRIOR APPLICATION NUMBER: US 60/074,725
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: US 60/096,409
PRIOR FILING DATE: 1998-08-13
NUMBER OF SEQ ID NOS: 28208
SEQ ID NO 15013
LENGTH: 620
TYPE: PRT
ORGANISM: Candida albicans
US-09-248-796A-15013

Query Match      85.4% Score 41; DB 2; Length 620;
Best Local Similarity 75.0%; Pred. No. 14;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      1 IHDIIEC 8
DB      493 LHDIIVLEC 500

RESULT 25
US-08-159-339A-573
Sequence 573, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Betteban
TITLE OF INVENTION: HLA Binding peptides and Their
TITLE OF INVENTION: Uses
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSER: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: PASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:

```

```

TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 573:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-573

Query Match      75.0% Score 36; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 1; 8;
Matches 7; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY      3 DIIEECV 9
DB      1 DIIEECV 7

RESULT 26
US-09-638-937-2
Sequence 2, Application US/09638937
Patent No. 6593514
GENERAL INFORMATION:
APPLICANT: Canoon, Edgar B
APPLICANT: Hitz, William D
APPLICANT: Ripp, Kevin G
TITLE OF INVENTION: METHOD FOR THE PRODUCTION OF CALENDIC ACID, AN UNUSUAL
TITLE OF INVENTION: FATY ACID CONTAINING DELTA-8,10,12 CONJUGATED DOUBLE
TITLE OF INVENTION: BONDS
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/09/638,937
CURRENT FILING DATE: 2000-08-15
PRIOR APPLICATION NUMBER: BB-1371-P1
PRIOR FILING DATE: 1999-08-16
NUMBER OF SEQ ID NOS: 25
SOFTWARE: Microsoft Office 97
SEQ ID NO 2
LENGTH: 374
TYPE: PRT
ORGANISM: Calendula officinalis
US-09-638-937-2

Query Match      75.0% Score 36; DB 2; Length 374;
Best Local Similarity 55.6%; Pred. No. 69;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 IHDIIECV 9
DB      47 LHDIIVTCI 55

RESULT 27
US-09-457-708-2
Sequence 2, Application US/09457708
Patent No. 6326483
GENERAL INFORMATION:
APPLICANT: Kwiatkowski, David J.
APPLICANT: Sampson, Julian R.
APPLICANT: Povey, Sue
APPLICANT: van Slegtenhorst, Marjon
APPLICANT: Halley, Dicky
TITLE OF INVENTION: Compositions and Methods Based U
TITLE OF INVENTION: Sclerosis-1 (TSC-1) Gene and Gene Product
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSER: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.

```

ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/457,708
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/42002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6585
TELEFAX: (202) 639-6604
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1164 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: Protein
HYPOTHEICAL: NO
ANTI-SENSE: NO
US-09-457-708-2

Query Match 75.0%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 2.2e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 HDIILEC 8
Db 253 HDVVIC 259

RESULT 28
US-09-950-046A-2
Sequence 2, Application US/09950046A
Patent No. 6548258
GENERAL INFORMATION:
APPLICANT: Kwiatkowski, David J.
Sampson, Julian R.
Povey, Sue
van Slegtenhore, Marjon
Halley, Dicky
TITLE OF INVENTION: Compositions and Methods Based Upon the Tubercous
Sclerosis-1 (TSC-1) Gene and Gene Product
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Vinson & Elkins
STREET: 1455 Pennsylvania Avenue, N.W.
CITY: Washington
STATE: D.C.
COUNTRY: U.S.
ZIP: 20004-1008
COMPUTER READABLE FORM:
MEDIUM TYPE: floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/950,046A
FILING DATE: 12-Sep-2001
CLASSIFICATION: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: Sanzo, Michael A.
REGISTRATION NUMBER: 36,912
REFERENCE/DOCKET NUMBER: BR1331/42002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 639-6585
TELEFAX: (202) 639-6604

INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 1164 amino acids
TYPE: amino acid
STRANDEDNESS: not relevant
TOPOLOGY: not relevant
MOLECULE TYPE: protein
HYPOTHEICAL: NO
ANTI-SENSE: NO
SEQUENCE DESCRIPTION: SEQ ID NO: 2:
US-09-950-046A-2

Query Match 75.0%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 2.2e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 HDIILEC 8
Db 253 HDVVIC 259

RESULT 29
US-09-976-594-989
Sequence 989, Application US/09976594
Patent No. 6673549
GENERAL INFORMATION:
APPLICANT: Furness, Michael
Buchbinder, Jenny
TITLE OF INVENTION: GENES EXPRESSED IN C3A LIVER CELL CULTURES TREATED WITH STEROIDS
FILE REFERENCE: PA-0041 US
CURRENT APPLICATION NUMBER: US/09/976,594
CURRENT FILING DATE: 2001-10-12
PRIOR APPLICATION NUMBER: 60/240,409
PRIOR FILING DATE: 2000-10-12
NUMBER OF SEQ ID NOS: 1143
SOFTWARE: PERL Program
SEQ ID NO 989
LENGTH: 1164
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
OTHER INFORMATION: Incyte ID No. 6673549 4215034CD1
US-09-976-594-989

Query Match 75.0%; Score 36; DB 2; Length 1164;
Best Local Similarity 57.1%; Pred. No. 2.2e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy 2 HDIILEC 8
Db 253 HDVVIC 259

RESULT 30
US-09-107-532A-5595
Sequence 5595, Application US/09107532A
Patent No. 6583275
GENERAL INFORMATION:
APPLICANT: Lynn A Doucette-Stamm and David Bush
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS
NUMBER OF SEQUENCES: 7310
CORRESPONDENCE ADDRESS:
ADDRESSEE: GENOME THERAPEUTICS CORPORATION
STREET: 100 Beaver Street
CITY: Waltham
STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD-ROM ISO9660
COMPUTER: PC

```

; OPERATING SYSTEM: <Unknown>
; SOFTWARE: ASCII
; CURRENT APPLICATION DATA:
;   APPLICATION NUMBER: US/09/107,532A
;   FILING DATE: 30-Jun-1998
; PRIOR APPLICATION DATA:
;   APPLICATION NUMBER: 60/085,598
;   FILING DATE: 14 May 1998
;   APPLICATION NUMBER: 60/051571
;   FILING DATE: July 2, 1997
; ATTORNEY/AGENT INFORMATION:
;   NAME: Arinello, Pamela Deneke
;   REGISTRATION NUMBER: 40,489
;   REFERENCE/DOCKET NUMBER: GTC-012
; TELECOMMUNICATION INFORMATION:
;   TELEPHONE: (781)893-5007
;   TELEFAX: (781)893-8277
; INFORMATION FOR SEQ ID NO: 5595:
;   SEQUENCE CHARACTERISTICS:
;     LENGTH: 86 amino acids
;     TYPE: amino acid
;     TOPOLOGY: linear
; MOLECULE TYPE: protein
; HYPOTHEICAL: YES
; ORIGINAL SOURCE:
;   ORGANISM: Enterococcus faecium
; FEATURE:
;   NAME/KEY: misc feature
;   LOCATION: (B) LOCATION 1...86
; SEQUENCE DESCRIPTION: SEQ ID NO: 5595:
US-09-107-532A-5595

Query Match          70.8%; Score 34; DB 2; Length 86;
Best Local Similarity 44.4%; Pred. No. 37;
Matches 4; Conservative 5; Mismatches 0; Indels 0; Gaps 0;

QY 1 HDIILCEV 9
   :|::|
DB 8 VYDLIVECV 16

RESULT 31
US-09-270-767-33608
; Sequence 33608, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
;   APPLICANT: Homburger et al.
;   TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
;   FILE REFERENCE: File Reference: 7326-094
;   CURRENT APPLICATION NUMBER: US/09/270,767
;   CURRENT FILING DATE: 1999-03-17
;   NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 33608
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-33608

Query Match          70.8%; Score 34; DB 2; Length 111;
Best Local Similarity 50.0%; Pred. No. 47;
Matches 4; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

QY 1 HDIILCE 8
   :|::|
DB 33 LHDILKIC 40

RESULT 32
US-09-270-767-48825
; Sequence 48825, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
```

```

; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 48825
; LENGTH: 111
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
US-09-270-767-48825

Query Match          70.8%; Score 34; DB 2; Length 111;
Best Local Similarity 50.0%; Pred. No. 47;
Matches 4; Conservative 4; Mismatches 0; Indels 0; Gaps 0;

QY 1 HDIILCE 8
   :|::|
DB 33 LHDILKIC 40

RESULT 33
US-09-489-039A-13381
; Sequence 13381, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
;   APPLICANT: Gary Breton et. al
;   TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
;   TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
;   FILE REFERENCE: 2709,2004001
;   CURRENT APPLICATION NUMBER: US/09/489,039A
;   CURRENT FILING DATE: 2000-01-27
;   PRIOR APPLICATION NUMBER: US 60/117,747
;   PRIOR FILING DATE: 1999-01-29
;   NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 13381
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-13381

Query Match          70.8%; Score 34; DB 2; Length 266;
Best Local Similarity 57.1%; Pred. No. 1.1e+02;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 HDIILCE 8
   :|::|
DB 138 HDLVDC 144

RESULT 34
US-09-638-937-4
; Sequence 4, Application US/09638937
; Patent No. 6593514
; GENERAL INFORMATION:
;   APPLICANT: Cahoon, Edgar B
;   APPLICANT: Hitz, William D
;   APPLICANT: Ripp, Kevin G
;   TITLE OF INVENTION: METHOD FOR THE PRODUCTION OF CALENDIC ACID, AN UNUSUAL
;   TITLE OF INVENTION: FATTY ACID CONTAINING DELTA-8,10,12 CONJUGATED DOUBLE
;   TITLE OF INVENTION: BONDS
;   FILE REFERENCE:
;   CURRENT APPLICATION NUMBER: US/09/638,937
;   CURRENT FILING DATE: 2000-08-15
;   PRIOR APPLICATION NUMBER: BB-1371-P1
;   PRIOR FILING DATE: 1999-08-16
;   NUMBER OF SEQ ID NOS: 25
; SOFTWARE: Microsoft Office 97
; SEQ ID NO 4
; LENGTH: 372
; TYPE: PRT
; ORGANISM: Calendula officinalis
```

US-09-638-937-4

Query Match 70.8%; Score 34; DB 2; Length 372;
Best Local Similarity 62.5%; Pred. No. 1.6e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Oy 2 HDIILECV 9
:|:|:|:|:
Db 48 HDIIVTCT 55

RESULT 35

US-09-107-532A-6713

Sequence 6713, Application US/09107532A

Patent No. 6583275

GENERAL INFORMATION:

APPLICANT: Lynn A Doucette-Stamm and David Bush

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS

NUMBER OF SEQUENCES: 7310

CORRESPONDENCE ADDRESS:

ADDRESS: GENOME THERAPEUTICS CORPORATION

STREET: 100 Beaver Street

CITY: Waltham

STATE: Massachusetts

COUNTRY: USA

ZIP: 02354

COMPUTER READABLE FORM:

MEDIUM TYPE: CD-ROM ISO9660

COMPUTER: PC

OPERATING SYSTEM: <Unknown>

SOFTWARE: ASCII

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/107.532A

FILING DATE: 30-Jun-1998

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 60/085,598

FILING DATE: 14 May 1998

APPLICATION NUMBER: 60/051571

FILING DATE: July 2, 1997

ATTORNEY/AGENT INFORMATION:

NAME: Arinello, Pamela Deneke

REGISTRATION NUMBER: 40,489

REFERENCE/DOCKET NUMBER: GTC-012

TELECOMMUNICATION INFORMATION:

TELEPHONE: (781)893-5007

TELEFAX: (781)893-8277

INFORMATION FOR SEQ ID NO: 6713:

SEQUENCE CHARACTERISTICS:

LENGTH: 426 amino acids

TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

HYPOTHEICAL: YES

ORIGINAL SOURCE:

ORGANISM: Enterococcus faecium

FEATURE:

NAME/KEY: misc feature

LOCATION: (B) LOCATION 1...426

SEQUENCE DESCRIPTION: SEQ ID NO: 6713:

US-09-107-532A-6713

Query Match 70.8%; Score 34; DB 2; Length 426;
Best Local Similarity 44.4%; Pred. No. 1.8e+02;
Matches 4; Conservative 5; Mismatches 0; Indels 0; Gaps 0;

Oy 1 IHDIILECV 9
:|:|:|:|:
Db 225 VYDLIVCEV 233

RESULT 36

US-09-252-991A-17124

Sequence 17124, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252.991A

CURRENT FILING DATE: 1999-02-18

PRIOR FILING DATE: 1998-02-18

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 17124

LENGTH: 956

TYPE: PRT

ORGANISM: Pseudomonas aeruginosa

US-09-252-991A-17124

Query Match 70.8%; Score 34; DB 2; Length 956;
Best Local Similarity 55.6%; Pred. No. 4.1e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Oy 1 IHDIILECV 9
:|:|:|:|:
Db 744 VADIVLECV 752

RESULT 37

US-09-252-991A-24727

Sequence 24727, Application US/09252991A

Patent No. 6551795

GENERAL INFORMATION:

APPLICANT: Marc J. Rubenfield et al.

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS

AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS

FILE REFERENCE: 107196.136

CURRENT APPLICATION NUMBER: US/09/252.991A

CURRENT FILING DATE: 1999-02-18

PRIOR FILING DATE: 1998-02-18

PRIOR APPLICATION NUMBER: US 60/094,190

PRIOR FILING DATE: 1998-07-27

NUMBER OF SEQ ID NOS: 33142

SEQ ID NO 24727

LENGTH: 295

TYPE: PRT

ORGANISM: Pseudomonas aeruginosa

US-09-252-991A-24727

Query Match 68.8%; Score 33; DB 2; Length 295;
Best Local Similarity 44.4%; Pred. No. 1.9e+02;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Oy 1 IHDIILECV 9
:|:|:|:|:
Db 264 IHDVLEAI 272

RESULT 38

US-09-107-532A-5486

Sequence 5486, Application US/09107532A

Patent No. 6583275

GENERAL INFORMATION:

APPLICANT: Lynn A Doucette-Stamm and David Bush

TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO

ENTEROCOCCUS FAECIUM FOR DIAGNOSTICS AND THERAPEUTICS

NUMBER OF SEQUENCES: 7310

CORRESPONDENCE ADDRESS:

ADDRESS: GENOME THERAPEUTICS CORPORATION

STREET: 100 Beaver Street

CITY: Waltham

```

STATE: Massachusetts
COUNTRY: USA
ZIP: 02354
COMPUTER READABLE FORM:
MEDIUM TYPE: CD/ROM ISO9660
OPERATING SYSTEM: <Unknown>
SOFTWARE: ASCII
CURRENT APPLICATION DATA:
  APPLICATION NUMBER: US/09/107,532A
  FILING DATE: 30-Jun-1998
PRIOR APPLICATION DATA:
  APPLICATION NUMBER: 60/085,598
  FILING DATE: 14 May 1998
  APPLICATION NUMBER: 60/051571
  FILING DATE: July 2, 1997
ATTORNEY/AGENT INFORMATION:
  NAME: Atinello, Pamela Deneke
  REGISTRATION NUMBER: 40,489
  REFERENCE/DOCKET NUMBER: GTC-012
TELECOMMUNICATION INFORMATION:
  TELEPHONE: (781)893-5007
  TELEFAX: (781)893-8277
INFORMATION FOR SEQ ID NO: 5486:
  SEQUENCE CHARACTERISTICS:
    LENGTH: 305 amino acids
    TYPE: amino acid
    TOPOLOGY: linear
MOLECULAR TYPE: protein
HYPOTHETICAL: YES
ORIGINAL SOURCE:
  ORGANISM: Enterococcus faecium
FEATURE:
  NAME/KEY: misc feature
  LOCATION: (8) LOCATION 1...305
  SEQUENCE DESCRIPTION: SEQ ID NO: 5486:
US-09-107-532A-5486

```

```

Query Match      68.8%; Score 33; DB 2; Length 305;
Best Local Similarity 85.7%; Pred. No. 2e+02;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

```

```

QY      3 DIIIECV 9
      |||||
      129 DIIIECM 135

```

```

RESULT 39
US-09-513-999C-5568
; Sequence 5568, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Mline Edwards, J.B.
; APPLICANT: Duclet, A.
; APPLICANT: Giordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 6783961
; FILE REFERENCE: 59 US2, REG
; CURRENT APPLICATION NUMBER: US/09/513,999C
; CURRENT FILING DATE: 2000-02-24
; PRIOR APPLICATION NUMBER: US 60/122,487
; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 36681
; SOFTWARE: Patent.pm
; SEQ ID NO 5568
; LENGTH: 119
; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: 39
; OTHER INFORMATION: Xaa=Asn or Thr
; FEATURE:

```

```

; NAME/KEY: UNSURE
; LOCATION: 113
; OTHER INFORMATION: Xaa= * or Trp
; US-09-513-999C-5568

```

```

Query Match      66.7%; Score 32; DB 2; Length 119;
Best Local Similarity 62.5%; Pred. No. 1.2e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      2 HDIIIECV 9
      |||||
      19 HNIIVECV 26

```

```

RESULT 40
US-09-270-767-60201
; Sequence 60201, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 60201
; LENGTH: 217
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-60201

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Query Match      66.7%; Score 32; DB 2; Length 217;
Best Local Similarity 75.0%; Pred. No. 2.2e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

```

```

QY      1 IHDIIECV 8
      |||||
      52 IHDIIECV 59

```

```

RESULT 41
US-09-252-991A-29181
; Sequence 29181, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 29181
; LENGTH: 227
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
; US-09-252-991A-29181

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Query Match      66.7%; Score 32; DB 2; Length 227;
Best Local Similarity 62.5%; Pred. No. 2.3e+02;
Matches 5; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

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QY      1 IHDIIECV 8
      |||||
      140 IHDFVNEC 147

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RESULT 42
US-09-328-352-8022
; Sequence 8022, Application US/09328352
; Patent No. 6562958
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
; FILE REFERENCE: GTC99-03PA
; CURRENT APPLICATION NUMBER: US/09/328,352
; CURRENT FILING DATE: 1999-06-04
; NUMBER OF SEQ ID NOS: 8252
; SEQ ID NO 8022
; LENGTH: 269
; TYPE: PRT
; ORGANISM: Acinetobacter baumannii
US-09-328-352-8022

Query Match      66.7%; Score 32; DB 2; Length 269;
Best Local Similarity 44.4%; Pred. No. 2.7e+02;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Cy 1 IHDIIECV 9
Db 137 VPDIVLDCI 145

RESULT 43
US-09-248-796A-22434
; Sequence 22434, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 22434
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Candida albicans
US-09-248-796A-22434

Query Match      66.7%; Score 32; DB 2; Length 343;
Best Local Similarity 100.0%; Pred. No. 3.4e+02;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 3 DIILIEC 8
Db 122 DIILIEC 127

RESULT 44
US-09-270-767-44745
; Sequence 44745, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-084
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patent In Ver. 2.0
; SEQ ID NO 44745
; LENGTH: 361
```

```
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-44745

Query Match      66.7%; Score 32; DB 2; Length 361;
Best Local Similarity 75.0%; Pred. No. 3.6e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Cy 1 IHDIIEC 8
Db 196 IHDIIEC 203

RESULT 45
US-08-426-509A-12
; Sequence 12, Application US/08426509A
; Patent No. 6326469
; GENERAL INFORMATION:
; APPLICANT: Ulrich, Axel
; APPLICANT: Gienhizsky, Mikhail
; TITLE OF INVENTION: NOVEL MEGAKARYOCYTIC PROTEIN
; TITLE OF INVENTION: TYROSINE KINASES
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York,
; STATE: NY
; COUNTRY: USA
; ZIP: 10036-2711
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: FastSeq Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/426,509A
; FILING DATE: 21-APR-1995
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 08/232,545
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: Coruzzi, Laura A
; REGISTRATION NUMBER: 30,742
; REFERENCE/DOCKET NUMBER: 7683-0074-999
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 212-790-9090
; TELEFAX: 212-869-9741
; TELEX: 66141 PENNIB
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 536 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: No. 6326469e
US-08-426-509A-12

Query Match      66.7%; Score 32; DB 2; Length 536;
Best Local Similarity 37.5%; Pred. No. 5.4e+02;
Matches 3; Conservative 5; Mismatches 0; Indels 0; Gaps 0;

Cy 1 IHDIIEC 8
Db 494 LHDVWQC 501

RESULT 46
US-08-232-545-12
```



```
; Sequence 12, Application US/08232545
; Patent No. 6506578
; GENERAL INFORMATION:
; APPLICANT: Ullrich, Axel
; APPLICANT: Gishizky, Mikhail
; APPLICANT: Sures, Irmann G.
; TITLE OF INVENTION: No. 6506578e1 Megakaryocytic Protein Tyrosine
; TITLE OF INVENTION: Kinases
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/232,545
; FILING DATE: 22-APR-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Coruzzi, Laura A.
; REGISTRATION NUMBER: 30,742
; REFERENCE/DOCKET NUMBER: 7683-050
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212)790-9090
; TELEFAX: (212)869-9741
; TELER: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 536 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
; US-08-232-545-12

Query Match      66.7%; Score 32; DB 2; Length 536;
Best Local Similarity 37.5%; Pred. No. 5.4e+02;
Matches 3; Conservative 5; Mismatches 0; Indels 0; Gaps 0;
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```
Cy 1 IHDIIIBC 8
Db 494 LHDVWVC 501
```

```
RESULT 47
US-09-977-261-12
; Sequence 12, Application US/09977261
; Patent No. 6908984
; GENERAL INFORMATION:
; APPLICANT: ULLRICH, AXEL
; APPLICANT: GISHIZKY, MIKHAIL
; APPLICANT: SURES, IRMANN G.
; TITLE OF INVENTION: NOVEL MEGAKARYOCYTIC PROTEIN TYROSINE KINASES
; FILE REFERENCE: 038602/1259
; CURRENT APPLICATION NUMBER: US/09/977,261
; CURRENT FILING DATE: 2001-10-16
; PRIOR APPLICATION NUMBER: 08/232,545
; PRIOR FILING DATE: 1994-04-22
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 12
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Gallus gallus
; US-09-977-261-12
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```
Query Match      66.7%; Score 32; DB 2; Length 536;
Best Local Similarity 37.5%; Pred. No. 5.4e+02;
Matches 3; Conservative 5; Mismatches 0; Indels 0; Gaps 0;
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```
Cy 1 IHDIIIBC 8
Db 494 LHDVWVC 501
```

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RESULT 48
PCT-US95-05008-12
; Sequence 12, Application PC/TUS9505008
; GENERAL INFORMATION:
; APPLICANT: Sugen, Inc.
; APPLICANT: 515 Galveston Drive
; APPLICANT: Redwood City, California 94063-4720
; APPLICANT: United States of America
; APPLICANT: Wiesenschaften E.V.
; APPLICANT: Hofgarten Str. 2
; APPLICANT: Munchen 80539
; APPLICANT: Germany
; TITLE OF INVENTION: Novel Megakaryocytic Protein Tyrosine
; TITLE OF INVENTION: Kinases
; NUMBER OF SEQUENCES: 21
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Pennie & Edmonds
; STREET: 1155 Avenue of the Americas
; CITY: New York
; STATE: New York
; COUNTRY: U.S.A.
; ZIP: 10036
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; OPERATING SYSTEM: IBM PC compatible
; SOFTWARE: PatentIn Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: PCT/US95/05008
; FILING DATE: 24-APR-1995
; CLASSIFICATION:
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 08/232,545
; FILING DATE: 22-APR-1994
; CLASSIFICATION:
; ATTORNEY/AGENT INFORMATION:
; NAME: Coruzzi, Laura A.
; REGISTRATION NUMBER: 30,742
; REFERENCE/DOCKET NUMBER: 7683-074
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (212)790-9090
; TELEFAX: (212)869-9741
; TELER: 66141 PENNIE
; INFORMATION FOR SEQ ID NO: 12:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 536 amino acids
; TYPE: amino acid
; STRANDEDNESS: unknown
; TOPOLOGY: unknown
; MOLECULE TYPE: protein
; PCT-US95-05008-12
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Query Match      66.7%; Score 32; DB 4; Length 536;
Best Local Similarity 37.5%; Pred. No. 5.4e+02;
Matches 3; Conservative 5; Mismatches 0; Indels 0; Gaps 0;
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Cy 1 IHDIIIBC 8
Db 494 LHDVWVC 501
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RESULT 49
US-09-270-767-45410
; Sequence 45410, Application US/09270767
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Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 45410
LENGTH: 584
TYPE: PRT
ORGANISM: Drosophila melanogaster
US-09-270-767-45410

Query Match      66.7%; Score 32; DB 2; Length 584;
Best Local Similarity 55.6%; Pred. No. 5.9e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 DHDILBCV 9
Db      172 LHDDILDCM 180

RESULT 50
US-09-907-794A-339
Sequence 339; Application US/09907794A
Patent No. 6635468
GENERAL INFORMATION:
APPLICANT: Genentech, Inc.
APPLICANT: Ashkenazi, Avi
APPLICANT: Botstein, David
APPLICANT: Desnoyers, Luc
APPLICANT: Balon, Dan L.
APPLICANT: Ferrara, Napoleone
APPLICANT: Filvaroff, Ellen
APPLICANT: Fong, Sherman
APPLICANT: Gao, Wei-Qiang
APPLICANT: Gerber, Hanspeter
APPLICANT: Gertschen, Mary E.
APPLICANT: Goddard, A.
APPLICANT: Godowski, Paul J.
APPLICANT: Grimaldi, Christopher J.
APPLICANT: Gurney, Austin L.
APPLICANT: Hillan, Kenneth, J.
APPLICANT: Kljavin, Ivar J.
APPLICANT: Mather, Jennie P.
APPLICANT: Pan, James
APPLICANT: Paoni, Nicholas P.
APPLICANT: Roy, Margaret Ann
APPLICANT: Stewart, Timothy A.
APPLICANT: Tumas, Daniel
APPLICANT: Williams, P. Mickey
APPLICANT: Wood, William, I.
TITLE OF INVENTION: Secreted and Transmembrane Polypeptides and Nucleic
FILE REFERENCE: 10466-14
CURRENT APPLICATION NUMBER: US/09/907,794A
CURRENT FILING DATE: 2001-07-17
PRIOR APPLICATION NUMBER: PCT/US00/04414
PRIOR FILING DATE: 2000-02-22
PRIOR APPLICATION NUMBER: US 60/143,048
PRIOR FILING DATE: 1999-07-07
PRIOR APPLICATION NUMBER: US 60/145,698
PRIOR FILING DATE: 1999-07-26
PRIOR APPLICATION NUMBER: US 60/146,222
PRIOR FILING DATE: 1999-07-28
PRIOR APPLICATION NUMBER: PCT/US99/20594
PRIOR FILING DATE: 1999-09-08
PRIOR APPLICATION NUMBER: PCT/US99/20944
PRIOR FILING DATE: 1999-09-13
PRIOR APPLICATION NUMBER: PCT/US99/21090
PRIOR FILING DATE: 1999-09-15
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PRIOR APPLICATION NUMBER: PCT/US99/21547
PRIOR FILING DATE: 1999-09-15
PRIOR APPLICATION NUMBER: PCT/US99/23089
PRIOR FILING DATE: 1999-10-05
PRIOR APPLICATION NUMBER: PCT/US99/28214
PRIOR FILING DATE: 1999-11-29
PRIOR APPLICATION NUMBER: PCT/US99/28313
PRIOR FILING DATE: 1999-11-30
PRIOR APPLICATION NUMBER: PCT/US99/28564
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/28565
PRIOR FILING DATE: 1999-12-02
PRIOR APPLICATION NUMBER: PCT/US99/30095
PRIOR FILING DATE: 1999-12-16
PRIOR APPLICATION NUMBER: PCT/US99/30911
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US99/30999
PRIOR FILING DATE: 1999-12-20
PRIOR APPLICATION NUMBER: PCT/US00/00219
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 423
SEQ ID NO 339
LENGTH: 772
TYPE: PRT
ORGANISM: Homo Sapien
US-09-907-794A-339

Query Match      66.7%; Score 32; DB 2; Length 772;
Best Local Similarity 71.4%; Pred. No. 7.8e+02;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY      3 DILBCV 9
Db      447 DILBCV 453
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Search completed: May 5, 2006, 04:48:29
job time : 22.7 secs

GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:18:14 ; Search time 56 Seconds
(without alignments)
67.151 Million cell updates/sec

Title: US-08-170-344-8
Perfect score: 48
Sequence: 1 IHDIILRCV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues

Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :
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2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep.*
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4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.pep.*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	48	100.0	10	4	US-10-128-711-72 Sequence 72, Appl
2	48	100.0	10	4	US-10-133-210-280 Sequence 22, Appl
3	48	100.0	15	4	US-10-476-570-22 Sequence 23, Appl
4	48	100.0	15	4	US-10-476-570-23 Sequence 10, Appl
5	48	100.0	21	4	US-10-476-570-10 Sequence 53, Appl
6	48	100.0	30	4	US-10-476-570-53 Sequence 4, Appl
7	48	100.0	30	5	US-10-858-384-4 Sequence 9, Appl
8	48	100.0	32	4	US-10-476-570-9 Sequence 19, Appl
9	48	100.0	33	4	US-10-177-390-6 Sequence 20, Appl
10	48	100.0	151	5	US-10-484-063-20 Sequence 27, Appl
11	48	100.0	151	5	US-10-484-063-27 Sequence 2, Appl
12	48	100.0	158	5	US-10-858-384-2 Sequence 16, Appl
13	48	100.0	158	5	US-10-367-057-16 Sequence 13, Appl
14	48	100.0	158	6	US-11-021-949-13 Sequence 2, Appl
15	48	100.0	171	4	US-10-472-724-2 Sequence 1, Appl
16	48	100.0	171	4	US-11-072-288-1 Sequence 1, Appl
17	48	100.0	243	6	US-09-367-309A-1 Sequence 4, Appl
18	48	100.0	266	4	US-10-000-903-4 Sequence 10, Appl
19	48	100.0	273	5	US-10-899-771-4 Sequence 6, Appl
20	48	100.0	273	5	US-10-000-903-10 Sequence 10, Appl
21	48	100.0	292	5	US-10-899-771-10 Sequence 6, Appl
22	48	100.0	371	5	US-10-000-903-6 Sequence 6, Appl
23	48	100.0	371	5	US-10-899-771-6 Sequence 14, Appl
24	48	100.0	390	5	US-10-000-903-14 Sequence 14, Appl
25	48	100.0	390	5	US-10-899-771-14 Sequence 10, Appl
26	48	100.0	536	4	US-10-367-095-10 Sequence 10, Appl
27	48	100.0	536	4	US-10-367-095-10 Sequence 10, Appl

28	48	100.0	536	4	US-10-368-046-10 Sequence 10, Appl
29	48	100.0	536	4	US-10-367-367-10 Sequence 10, Appl
30	48	100.0	536	5	US-10-918-337-10 Sequence 547, Appl
31	44	91.7	9	4	US-10-777-053-547 Sequence 547, Appl
32	44	91.7	9	4	US-10-837-217-547 Sequence 24, Appl
33	44	91.7	15	4	US-10-476-570-24 Sequence 16507, A
34	39	81.2	297	4	US-10-424-599-16507 Sequence 275411, A
35	39	81.2	298	4	US-10-424-599-16507 Sequence 193535, A
36	38	79.2	148	6	US-10-425-115-193535 Sequence 10846, A
37	38	79.2	554	5	US-10-739-930-10846 Sequence 125430, A
38	37	77.1	58	4	US-10-437-963-125430 Sequence 219049, A
39	37	77.1	58	4	US-10-424-599-219049 Sequence 33593, A
40	36	75.0	70	4	US-10-029-386-33593 Sequence 205547, A
41	36	75.0	83	4	US-10-425-115-205547 Sequence 351, Appl
42	36	75.0	158	6	US-11-021-949-361 Sequence 58813, A
43	36	75.0	274	4	US-10-425-114-58813 Sequence 65704, A
44	36	75.0	274	4	US-10-425-114-65704 Sequence 145297, A
45	36	75.0	291	4	US-10-437-963-145297 Sequence 187372, A
46	36	75.0	294	4	US-10-424-599-187372 Sequence 52907, A
47	36	75.0	294	4	US-10-425-114-52907 Sequence 36891, A
48	36	75.0	297	4	US-10-767-701-36891 Sequence 345578, A
49	36	75.0	304	4	US-10-425-115-345578 Sequence 146120, A
50	36	75.0	306	4	US-10-437-963-126120 Sequence 149433, A
51	36	75.0	306	4	US-10-437-963-149433 Sequence 50748, A
52	36	75.0	312	4	US-10-425-114-50748 Sequence 151133, A
53	36	75.0	356	4	US-10-437-963-151133 Sequence 2, Appl
54	36	75.0	374	3	US-09-852-399-2 Sequence 4880, Appl
55	36	75.0	374	3	US-10-464-631-2 Sequence 15537, A
56	36	75.0	374	5	US-10-732-923-4880 Sequence 96, Appl
57	36	75.0	503	4	US-10-437-963-153537 Sequence 13152, A
58	36	75.0	789	4	US-10-342-844-86 Sequence 1152, A
59	36	75.0	1100	6	US-11-097-143-13152 Sequence 2, Appl
60	36	75.0	1164	3	US-09-950-046A-2 Sequence 793, Appl
61	36	75.0	1164	3	US-10-719-993-793 Sequence 794, Appl
62	36	75.0	1164	5	US-10-719-993-795 Sequence 795, Appl
63	36	75.0	1164	5	US-10-424-599-261856 Sequence 261856, A
64	36	75.0	108	4	US-10-437-963-135239 Sequence 5853, Appl
65	35	72.9	121	6	US-11-097-143-5853 Sequence 13044, A
66	35	72.9	124	6	US-11-097-143-21847 Sequence 21847, A
67	35	72.9	145	5	US-10-732-923-21847 Sequence 153797, A
68	35	72.9	150	4	US-10-437-963-153797 Sequence 31, Appl
69	35	72.9	162	6	US-11-021-949-31 Sequence 29716, A
70	35	72.9	271	4	US-10-767-701-39716 Sequence 27112, A
71	35	72.9	286	4	US-10-424-599-274112 Sequence 13557, A
72	35	72.9	334	5	US-10-368-493-13557 Sequence 20816, A
73	35	72.9	335	5	US-10-732-923-18204 Sequence 18204, A
74	35	72.9	336	4	US-10-425-115-262820 Sequence 115928, A
75	35	72.9	336	4	US-10-437-963-115928 Sequence 335998, A
76	35	72.9	570	4	US-10-425-115-359988 Sequence 194068, A
77	35	72.9	107	4	US-10-425-115-194068 Sequence 226149, A
78	34	70.8	117	4	US-10-425-115-226149 Sequence 6390, Appl
79	34	70.8	117	4	US-11-097-143-6390 Sequence 274871, A
80	34	70.8	123	6	US-10-425-115-274871 Sequence 335, Appl
81	34	70.8	148	6	US-11-021-949-359 Sequence 14374, A
82	34	70.8	149	6	US-10-767-701-43474 Sequence 535902, A
83	34	70.8	165	4	US-10-368-493-13677 Sequence 535902, A
84	34	70.8	241	4	US-10-282-122A-55972 Sequence 55972, A
85	34	70.8	246	4	US-10-282-122A-55972 Sequence 293913, A
86	34	70.8	251	4	US-10-282-122A-55972 Sequence 358748, A
87	34	70.8	265	4	US-10-425-115-358748 Sequence 292915, A
88	34	70.8	311	4	US-10-425-115-292915 Sequence 146645, A
89	34	70.8	315	4	US-10-437-963-146645 Sequence 77597, A
90	34	70.8	325	4	US-10-464-631-4 Sequence 4681, Appl
91	34	70.8	358	4	US-10-732-923-4881 Sequence 50390, A
92	34	70.8	372	5	US-10-437-963-199530 Sequence 150342, A
93	34	70.8	451	4	US-10-425-114-50342 Sequence 84, Appl
94	34	70.8	460	4	US-09-976-782-84 Sequence 13562, A
95	34	70.8	505	3	US-10-732-923-13562 Sequence 13562, A
96	34	70.8	505	5	US-10-732-923-13562 Sequence 13562, A

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102	34	70.8	627	4	US-10-425-115-274872	Sequence 274872, A	175	32	66.7	160	6	US-11-021-949-32	Sequence 32, Appl
103	34	70.8	648	6	US-11-097-143-13239	Sequence 13239, A	176	32	66.7	173	4	US-10-424-599-206501	Sequence 206501, A
104	34	70.8	665	6	US-11-097-143-19287	Sequence 19287, A	177	32	66.7	179	4	US-10-403-571-50	Sequence 50, Appl
105	34	70.8	774	4	US-10-437-963-135395	Sequence 135395, A	178	32	66.7	212	4	US-10-437-963-125445	Sequence 125445, A
106	34	70.8	787	4	US-10-282-122A-50697	Sequence 50697, A	179	32	66.7	217	4	US-10-437-963-137677	Sequence 137677, A
107	34	70.8	911	4	US-10-437-963-198474	Sequence 198474, A	180	32	66.7	218	4	US-10-424-599-141519	Sequence 141519, A
108	34	70.8	921	4	US-10-437-963-161130	Sequence 161130, A	181	32	66.7	245	4	US-10-437-963-184683	Sequence 184683, A
109	34	70.8	1269	4	US-10-437-963-190985	Sequence 190985, A	182	32	66.7	283	5	US-10-739-930-9600	Sequence 9600, Ap
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111	34	70.8	2737	5	US-10-450-763-37397	Sequence 37397, A	184	32	66.7	289	4	US-10-437-963-179146	Sequence 179146, A
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117	33	68.8	86	4	US-10-424-599-263346	Sequence 263346, A	190	32	66.7	334	4	US-10-425-115-31896	Sequence 317896, A
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121	33	68.8	138	4	US-10-767-701-34127	Sequence 34127, A	194	32	66.7	384	4	US-10-425-114-65198	Sequence 65198, A
122	33	68.8	148	6	US-11-021-949-17	Sequence 17, Appl	195	32	66.7	387	6	US-11-097-143-18264	Sequence 18264, A
123	33	68.8	149	6	US-11-021-949-16	Sequence 16, Appl	196	32	66.7	418	4	US-10-282-122A-49150	Sequence 49150, A
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125	33	68.8	157	3	US-09-833-245-110	Sequence 110, App	198	32	66.7	425	3	US-09-660-670-105	Sequence 105, App
126	33	68.8	169	3	US-09-938-803-2	Sequence 2, Appli	199	32	66.7	425	4	US-10-327-646-105	Sequence 105, App
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128	33	68.8	169	5	US-10-993-986-2	Sequence 2, Appl1	201	32	66.7	437	5	US-10-732-923-9879	Sequence 9879, Ap
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130	33	68.8	281	4	US-10-424-599-275162	Sequence 275162, A	203	32	66.7	447	5	US-10-732-923-20229	Sequence 20229, A
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133	33	68.8	302	4	US-10-424-599-176897	Sequence 176897, A	206	32	66.7	506	4	US-10-972-963-186	Sequence 186, App
134	33	68.8	310	4	US-10-368-493-32996	Sequence 32996, A	207	32	66.7	506	4	US-10-295-027-169	Sequence 169, App
135	33	68.8	313	6	US-11-097-143-21081	Sequence 21081, A	208	32	66.7	506	4	US-10-188-832-39	Sequence 39, Appl
136	33	68.8	334	4	US-10-424-599-188017	Sequence 188017, A	209	32	66.7	536	3	US-09-977-260-12	Sequence 12, Appl
137	33	68.8	347	4	US-10-282-122A-63007	Sequence 63007, A	210	32	66.7	536	5	US-09-977-261-11	Sequence 11, Appl
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142	33	68.8	916	4	US-10-437-963-167102	Sequence 167102, A	215	32	66.7	635	6	US-11-097-143-1554	Sequence 1554, Ap
143	33	68.8	996	4	US-10-437-963-187715	Sequence 187715, A	216	32	66.7	716	4	US-10-437-963-130939	Sequence 130939, A
144	33	68.8	1087	6	US-11-097-143-7296	Sequence 7296, Ap	217	32	66.7	743	4	US-10-387-226-368	Sequence 368, App
145	33	68.8	1118	4	US-10-437-963-192769	Sequence 192769, A	218	32	66.7	761	3	US-09-895-912X-10	Sequence 10, Appl
146	33	68.8	1248	4	US-10-437-963-126940	Sequence 126940, A	219	32	66.7	767	4	US-10-437-963-136065	Sequence 136065, A
147	33	68.8	1405	4	US-10-437-963-156135	Sequence 156135, A	220	32	66.7	772	3	US-09-909-088X-339	Sequence 339, App
148	33	68.8	1523	4	US-10-473-576-5	Sequence 5, Appl1	221	32	66.7	772	3	US-09-909-088X-339	Sequence 339, App
149	33	68.8	1729	4	US-10-437-963-177639	Sequence 177639, A	222	32	66.7	772	3	US-09-905-291X-339	Sequence 339, App
150	33	68.8	1830	4	US-10-437-963-189860	Sequence 189860, A	223	32	66.7	772	3	US-09-905-291X-339	Sequence 339, App
151	33	68.8	1839	4	US-10-032-585-7605	Sequence 7605, Ap	224	32	66.7	772	3	US-09-902-853-339	Sequence 339, App
152	33	68.8	2029	4	US-10-087-684-38	Sequence 38, Appl	225	32	66.7	772	3	US-09-907-824-339	Sequence 339, App
153	33	68.8	2037	4	US-10-218-779-39	Sequence 39, Appl	226	32	66.7	772	3	US-09-907-841-339	Sequence 339, App
154	33	68.8	2039	4	US-10-087-684-39	Sequence 39, Appl	227	32	66.7	772	3	US-09-904-011-339	Sequence 339, App
155	33	68.8	2037	4	US-10-218-779-39	Sequence 39, Appl	228	32	66.7	772	3	US-09-903-640-339	Sequence 339, App
156	33	68.8	2037	6	US-11-097-143-42576	Sequence 42576, A	229	32	66.7	772	3	US-09-908-093-339	Sequence 339, App
157	33	68.8	2037	4	US-10-437-963-125761	Sequence 125761, A	230	32	66.7	772	3	US-09-906-742-339	Sequence 339, App
158	33	66.7	53	3	US-09-864-761-46784	Sequence 46784, A	231	32	66.7	772	3	US-09-906-838-339	Sequence 339, App
159	32	66.7	36	3	US-10-424-599-195652	Sequence 195652, A	232	32	66.7	772	3	US-09-907-613-339	Sequence 339, App
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161	32	66.7	77	3	US-09-864-761-40148	Sequence 40148, A	234	32	66.7	772	3	US-09-904-859-339	Sequence 339, App
162	32	66.7	79	3	US-10-424-599-230323	Sequence 230323, A	235	32	66.7	772	3	US-09-909-204-339	Sequence 339, App
163	32	66.7	93	3	US-09-942-446-10	Sequence 10, Appl	236	32	66.7	772	3	US-09-904-820-339	Sequence 339, App
164	32	66.7	94	4	US-10-425-115-320993	Sequence 320993, A	237	32	66.7	772	3	US-09-904-886-339	Sequence 339, App
165	32	66.7	96	4	US-10-437-963-123433	Sequence 123433, A	238	32	66.7	772	3	US-09-906-646-339	Sequence 339, App
166	32	66.7	101	4	US-10-425-115-244198	Sequence 244198, A	239	32	66.7	772	3	US-09-906-700-339	Sequence 339, App
167	32	66.7	116	5	US-10-723-860-705	Sequence 705, App	240	32	66.7	772	3	US-09-903-786-339	Sequence 339, App
168	32	66.7	119	4	US-10-767-701-55749	Sequence 55749, A	241	32	66.7	772	3	US-09-902-903-339	Sequence 339, App
169	32	66.7	133	4	US-10-425-114-51081	Sequence 51081, A	242	32	66.7	772	3	US-09-903-749X-339	Sequence 339, App
170	32	66.7	140	3	US-09-764-891-3791	Sequence 3791, Ap	243	32	66.7	772	3	US-09-904-119-339	Sequence 339, App
171	32	66.7	144	3	US-10-425-115-296278	Sequence 296278, A	244	32	66.7	772	3	US-09-904-956-339	Sequence 339, App
172	32	66.7	149	6	US-11-021-949-15	Sequence 15, Appl	245	32	66.7	772	3	US-09-902-736-339	Sequence 339, App
173	32	66.7	155	3	US-09-864-761-35049	Sequence 35049, A	246	32	66.7	772	3	US-09-907-794-339	Sequence 339, App

247	32	66.7	772	3	US-09-903-943-339	Sequence 339, App	320	32	66.7	958	6	US-11-097-143-15084	Sequence 15084, A
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253	32	66.7	772	3	US-09-909-064-339	Sequence 339, App	326	32	66.7	1662	4	US-10-437-963-125137	Sequence 125137, A
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257	32	66.7	772	3	US-09-905-348-339	Sequence 339, App	330	32	66.7	2342	6	US-11-097-143-8544	Sequence 8544, Ap
258	32	66.7	772	3	US-09-905-088-339	Sequence 339, App	331	32	66.7	2443	6	US-11-097-143-8355	Sequence 8355, Ap
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262	32	66.7	772	3	US-09-902-634-339	Sequence 339, App	335	31.5	1011	4	US-10-437-963-134127	Sequence 134127, A	
263	32	66.7	772	3	US-09-902-713-339	Sequence 339, App	336	31	64.6	44	3	US-09-397-945-249	Sequence 249, App
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265	32	66.7	772	3	US-09-902-615-339	Sequence 339, App	338	31	64.6	46	4	US-10-425-115-286093	Sequence 286093, A
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267	32	66.7	772	3	US-09-906-760A-339	Sequence 339, App	340	31	64.6	48	4	US-10-001-876-195	Sequence 195, App
268	32	66.7	772	3	US-09-903-823-339	Sequence 339, App	341	31	64.6	49	4	US-10-282-122A-57355	Sequence 57355, A
269	32	66.7	772	3	US-09-907-652-339	Sequence 339, App	342	31	64.6	51	4	US-10-437-963-203708	Sequence 203708, A
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271	32	66.7	772	3	US-09-902-979-339	Sequence 339, App	344	31	64.6	52	3	US-09-984-429-111	Sequence 111, App
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273	32	66.7	772	3	US-09-906-815A-339	Sequence 339, App	346	31	64.6	53	4	US-10-437-963-125154	Sequence 125154, A
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286	32	66.7	772	3	US-09-907-728-339	Sequence 339, App	359	31	64.6	103	4	US-10-767-701-56094	Sequence 56094, A
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289	32	66.7	772	3	US-09-906-722A-339	Sequence 339, App	362	31	64.6	108	6	US-11-002-562-6	Sequence 6, App1
290	32	66.7	772	3	US-09-908-576-339	Sequence 339, App	363	31	64.6	110	4	US-10-282-122A-52718	Sequence 52718, A
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293	32	66.7	772	4	US-10-299-976-339	Sequence 339, App	366	31	64.6	116	4	US-10-282-122A-51739	Sequence 51739, A
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295	32	66.7	772	4	US-10-211-858-20	Sequence 20, App1	368	31	64.6	118	4	US-10-168-506-24	Sequence 24, App1
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298	32	66.7	772	4	US-10-449-656-339	Sequence 339, App	371	31	64.6	122	4	US-10-767-701-58356	Sequence 58356, A
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305	32	66.7	772	5	US-10-978-255-339	Sequence 339, App	378	31	64.6	164	4	US-10-424-599-147892	Sequence 147892, A
306	32	66.7	772	5	US-10-970-823-339	Sequence 339, App	379	31	64.6	168	4	US-10-425-115-193767	Sequence 193767, A
307	32	66.7	780	4	US-10-420-495-11	Sequence 11, App1	380	31	64.6	169	4	US-10-264-237-2283	Sequence 2283, App
308	32	66.7	780	4	US-10-437-963-174758	Sequence 174758, A	381	31	64.6	170	4	US-10-424-599-250124	Sequence 250124, A
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310	32	66.7	780	5	US-10-631-467-14	Sequence 667, App	383	31	64.6	172	4	US-10-767-701-32405	Sequence 32405, A
311	32	66.7	780	5	US-10-631-467-667	Sequence 1436, App	384	31	64.6	173	4	US-10-425-115-358724	Sequence 358724, A
312	32	66.7	780	5	US-10-631-467-1456	Sequence 930, App	385	31	64.6	174	5	US-10-947-979-18	Sequence 18, App1
313	32	66.7	790	4	US-10-295-027-930	Sequence 1582, Ap	386	31	64.6	176	4	US-10-425-115-236213	Sequence 236213, A
314	32	66.7	821	4	US-10-408-765A-1582	Sequence 173301, A	387	31	64.6	176	5	US-10-947-979-10	Sequence 10, App1
315	32	66.7	857	4	US-10-437-963-173301	Sequence 184762, A	388	31	64.6	178	4	US-10-106-658-6566	Sequence 6566, Ap
316	32	66.7	870	4	US-10-437-963-184762	Sequence 1736075, A	389	31	64.6	181	4	US-10-425-115-229266	Sequence 229266, A
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318	32	66.7	938	4	US-10-437-963-173296	Sequence 173296, A	391	31	64.6	184	5	US-10-947-979-52	Sequence 52, App1
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395	31	64.6	230	4	US-10-767-701-42299	Sequence 42299, A	468	31	64.6	554	5	US-10-450-763-18188	Sequence 18188, A
396	31	64.6	232	5	US-10-501-282-552	Sequence 552, App	469	31	64.6	559	4	US-10-424-599-288466	Sequence 288466, A
397	31	64.6	235	5	US-10-777-144-13	Sequence 13, Appl	470	31	64.6	592	4	US-10-425-114-45837	Sequence 45837, A
398	31	64.6	235	5	US-10-777-145-13	Sequence 13, Appl	471	31	64.6	604	5	US-10-732-923-98877	Sequence 98877, Ap
399	31	64.6	235	5	US-10-777-186-13	Sequence 13, Appl	472	31	64.6	607	6	US-11-002-562-22	Sequence 22, Appl
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406	31	64.6	275	4	US-10-021-660-98	Sequence 98, Appl	479	31	64.6	690	6	US-10-044-543-6	Sequence 6, Appl
407	31	64.6	275	4	US-10-211-462-103	Sequence 103, App	480	31	64.6	690	6	US-10-021-664-2	Sequence 2, Appl
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459	31	64.6	491	5	US-10-481-032A-594	Sequence 594, App	532	31	64.6	1948	4	US-10-425-115-344401	Sequence 344401, A
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464	31	64.6	545	4	US-10-437-963-125450	Sequence 125450, A	537	31	64.6	1948	4	US-10-425-115-344401	Sequence 344401, A
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543	30	62.5	46	3	US-09-820-649-345	Sequence 345, App	616	30	62.5	294	6	US-09-883-249-1	Sequence 1, App1
544	30	62.5	46	4	US-10-160-162-345	Sequence 345, App	617	30	62.5	296	3	US-11-097-143-35331	Sequence 35331, A
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588	30	62.5	179	4	US-10-424-599-242161	Sequence 242161, A	661	30	62.5	425	4	US-10-369-493-7880	Sequence 7880, App
589	30	62.5	180	6	US-10-425-114-45820	Sequence 45820, A	662	30	62.5	427	4	US-10-425-114-65961	Sequence 65961, A
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595	30	62.5	213	4	US-10-024-579-2	Sequence 2, App1	668	30	62.5	445	5	US-10-619-533-23	Sequence 23, App1
596	30	62.5	213	4	US-10-168-651-11	Sequence 11, App1	669	30	62.5	446	4	US-10-282-122A-53394	Sequence 53394, A
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598	30	62.5	244	4	US-10-042-865-85	Sequence 85, App1	671	30	62.5	486	4	US-10-424-599-191194	Sequence 191194, A
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697	30	62.5	623	4	US-10-296-115-1190	Sequence 1190, Ap	770	30	62.5	945	4	US-10-131-824A-146	Sequence 146, App
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702	30	62.5	635	4	US-10-104-047-2449	Sequence 2449, Ap	775	30	62.5	945	4	US-10-147-517-146	Sequence 146, App
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706	30	62.5	756	4	US-10-437-963-194428	Sequence 194428, A	779	30	62.5	945	4	US-10-147-527-146	Sequence 146, App
707	30	62.5	757	6	US-11-097-143-25935	Sequence 25935, A	780	30	62.5	945	4	US-10-121-041-146	Sequence 146, App
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724	30	62.5	933	4	US-10-218-779-4	Sequence 4, Appl	797	30	62.5	945	4	US-10-125-926A-146	Sequence 146, App
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867	30	62.5	945	4	US-10-123-912-146	Sequence 146, App	940	30	62.5	945	4	US-10-142-887-146	Sequence 146, App
868	30	62.5	945	4	US-10-192-007-146	Sequence 146, App	941	30	62.5	945	4	US-10-142-888-146	Sequence 146, App
869	30	62.5	945	4	US-10-194-359-146	Sequence 146, App	942	30	62.5	945	4	US-10-143-034-146	Sequence 146, App
870	30	62.5	945	4	US-10-127-847A-146	Sequence 146, App	943	30	62.5	945	4	US-10-143-116-146	Sequence 146, App
871	30	62.5	945	4	US-10-137-866-146	Sequence 146, App	944	30	62.5	945	4	US-10-144-982-146	Sequence 146, App
872	30	62.5	945	4	US-10-146-726-146	Sequence 146, App	945	30	62.5	945	4	US-10-145-015-146	Sequence 146, App
873	30	62.5	945	4	US-10-146-727-146	Sequence 146, App	946	30	62.5	945	4	US-10-145-090-146	Sequence 146, App
874	30	62.5	945	4	US-10-146-788-146	Sequence 146, App	947	30	62.5	945	4	US-10-145-091-146	Sequence 146, App
875	30	62.5	945	4	US-10-152-380-146	Sequence 146, App	948	30	62.5	945	4	US-10-145-629-146	Sequence 146, App
876	30	62.5	945	4	US-10-153-934-146	Sequence 146, App	949	30	62.5	945	4	US-10-145-629-146	Sequence 146, App
877	30	62.5	945	4	US-10-140-807-146	Sequence 146, App	950	30	62.5	945	4	US-10-145-630-146	Sequence 146, App
878	30	62.5	945	4	US-10-140-924-146	Sequence 146, App	951	30	62.5	945	4	US-10-145-737-146	Sequence 146, App
879	30	62.5	945	4	US-10-140-924-146	Sequence 146, App	952	30	62.5	945	4	US-10-145-752-146	Sequence 146, App
880	30	62.5	945	4	US-10-141-698-146	Sequence 146, App	953	30	62.5	945	4	US-10-145-754-146	Sequence 146, App
881	30	62.5	945	4	US-10-141-702-146	Sequence 146, App	954	30	62.5	945	4	US-10-145-755-146	Sequence 146, App
882	30	62.5	945	4	US-10-141-704-146	Sequence 146, App	955	30	62.5	945	4	US-10-145-818-146	Sequence 146, App
883	30	62.5	945	4	US-10-142-421-146	Sequence 146, App	956	30	62.5	945	4	US-10-145-820-146	Sequence 146, App
884	30	62.5	945	4	US-10-142-432-146	Sequence 146, App	957	30	62.5	945	4	US-10-145-872-146	Sequence 146, App
885	30	62.5	945	4	US-10-142-767-146	Sequence 146, App	958	30	62.5	945	4	US-10-145-873-146	Sequence 146, App
886	30	62.5	945	4	US-10-143-033-146	Sequence 146, App	959	30	62.5	945	4	US-10-147-481-146	Sequence 146, App
887	30	62.5	945	4	US-10-144-994-146	Sequence 146, App	960	30	62.5	945	4	US-10-147-482-146	Sequence 146, App
888	30	62.5	945	4	US-10-145-628-146	Sequence 146, App	961	30	62.5	945	4	US-10-147-503-146	Sequence 146, App
889	30	62.5	945	4	US-10-145-746-146	Sequence 146, App	962	30	62.5	945	4	US-10-147-522-146	Sequence 146, App
890	30	62.5	945	4	US-10-145-748-146	Sequence 146, App	963	30	62.5	945	4	US-10-152-401-146	Sequence 146, App
891	30	62.5	945	4	US-10-145-823-146	Sequence 146, App	964	30	62.5	945	4	US-10-157-783-146	Sequence 146, App
892	30	62.5	945	4	US-10-145-826-146	Sequence 146, App	965	30	62.5	945	4	US-10-158-792-146	Sequence 146, App
893	30	62.5	945	4	US-10-145-826-146	Sequence 146, App	966	30	62.5	945	4	US-10-158-792-146	Sequence 146, App
894	30	62.5	945	4	US-10-145-876-146	Sequence 146, App	967	30	62.5	945	4	US-10-143-035-146	Sequence 146, App
895	30	62.5	945	4	US-10-145-959-146	Sequence 146, App	968	30	62.5	945	4	US-10-145-751-146	Sequence 146, App
896	30	62.5	945	4	US-10-146-724-146	Sequence 146, App	969	30	62.5	945	4	US-10-145-822-146	Sequence 146, App
897	30	62.5	945	4	US-10-146-725-146	Sequence 146, App	970	30	62.5	945	4	US-10-145-824-146	Sequence 146, App
898	30	62.5	945	4	US-10-146-795-146	Sequence 146, App	971	30	62.5	945	4	US-10-145-827-146	Sequence 146, App
899	30	62.5	945	4	US-10-147-495-146	Sequence 146, App	972	30	62.5	945	4	US-10-145-869-146	Sequence 146, App
900	30	62.5	945	4	US-10-147-501-146	Sequence 146, App	973	30	62.5	945	4	US-10-145-875-146	Sequence 146, App
901	30	62.5	945	4	US-10-147-504-146	Sequence 146, App	974	30	62.5	945	4	US-10-145-877-146	Sequence 146, App
902	30	62.5	945	4	US-10-147-506-146	Sequence 146, App	975	30	62.5	945	4	US-10-145-958-146	Sequence 146, App
903	30	62.5	945	4	US-10-147-509-146	Sequence 146, App	976	30	62.5	945	4	US-10-146-787-146	Sequence 146, App

Sequence	US-10-146-793-146	Sequence	146, App
977	30 62.5 945 4	Sequence 146, App	
978	30 62.5 945 4	Sequence 146, App	
979	30 62.5 945 4	Sequence 146, App	
980	30 62.5 945 4	Sequence 146, App	
981	30 62.5 945 4	Sequence 146, App	
982	30 62.5 945 4	Sequence 146, App	
983	30 62.5 945 4	Sequence 146, App	
984	30 62.5 945 4	Sequence 146, App	
985	30 62.5 945 4	Sequence 146, App	
986	30 62.5 945 4	Sequence 146, App	
987	30 62.5 945 4	Sequence 146, App	
988	30 62.5 945 4	Sequence 146, App	
989	30 62.5 945 4	Sequence 146, App	
990	30 62.5 945 4	Sequence 146, App	
991	30 62.5 945 4	Sequence 146, App	
992	30 62.5 945 4	Sequence 146, App	
993	30 62.5 945 4	Sequence 146, App	
994	30 62.5 945 4	Sequence 146, App	
995	30 62.5 945 4	Sequence 146, App	
996	30 62.5 945 4	Sequence 146, App	
997	30 62.5 945 4	Sequence 146, App	
998	30 62.5 945 4	Sequence 146, App	
999	30 62.5 945 4	Sequence 146, App	
1000	30 62.5 945 4	Sequence 146, App	

ALIGNMENTS

RESULT 1
US-10-128-711-72
; Sequence 72, Application US/10128711
; Publication No. US2003009634A1
GENERAL INFORMATION:
APPLICANT: VITIELLO, Maria A.
CHRSTNUT, Robert W.
SETTE, Alessandro D.
CELIS, Esteaban
GRAY, Howard
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ELICITING
CTL IMMUNITY
NUMBER OF SEQUENCES: 153
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend Kourie and Crew
STREET: Stewart Street Tower, One Market Plaza
CITY: San Francisco
STATE: California
COUNTRY: US
ZIP: 94105-1493
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/128,711
FILING DATE: 22-Apr-2002
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/197,484
FILING DATE: 16-FEB-1994
APPLICATION NUMBER: US 07/935,811
FILING DATE: 26-AUG-1992
APPLICATION NUMBER: US 07/874,491
FILING DATE: 27-APR-1992
APPLICATION NUMBER: US 07/827,682
FILING DATE: 29-JAN-1992
APPLICATION NUMBER: US 07/749,568
FILING DATE: 26-AUG-1991
ATTORNEY/AGENT INFORMATION:
NAME: Parmelee, Steven W
REGISTRATION NUMBER: 31,990
REFERENCE/DOCKET NUMBER: 14137-26-4

TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 467-9600
TELEFAX: (206) 623-6793
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: unknown
TOPOLOGY: unknown
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 72:
US-10-128-711-72

Query Match 100.0%; Score 48; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.034;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIIECV 9
Db 2 IHDIIECV 10

RESULT 2
US-10-133-210-280
; Sequence 280, Application US/10133210
; Publication No. US20030103964A1
GENERAL INFORMATION:
APPLICANT: Delisi, Charles
APPLICANT: Berzofsky, Jay
APPLICANT: Gulukota, Kamalakara
APPLICANT: Vaccaro, Dennis
APPLICANT: Wang, Zhiping
APPLICANT: Zhang, Chao
TITLE OF INVENTION: METHODS FOR DESIGNING MOLECULAR CONJUGATES AND
FILE REFERENCE: BU-035AX
CURRENT APPLICATION NUMBER: US/10/133,210
CURRENT FILING DATE: 2002-04-26
NUMBER OF SEQ ID NOS: 281
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 280
LENGTH: 10
TYPE: PPT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Synthetic
US-10-133-210-280

Query Match 100.0%; Score 48; DB 4; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.034;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIIECV 9
Db 2 IHDIIECV 10

RESULT 3
US-10-476-570-22
; Sequence 22, Application US/10476570
; Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLERS, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04

```
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 22
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 24-38
US-10-476-570-22
```

```
Query Match          100.0%; Score 48; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.051;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHHIILECV 9
Db 7 IHHIILECV 15
```

```
RESULT 4
US-10-476-570-23
; Sequence 23, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 23
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 28-42
US-10-476-570-23
```

```
Query Match          100.0%; Score 48; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.051;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHHIILECV 9
Db 3 IHHIILECV 11
```

```
RESULT 5
US-10-476-570-10
; Sequence 10, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
```

```
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 10
; LENGTH: 21
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 30-50
US-10-476-570-10
```

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Query Match          100.0%; Score 48; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.073;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
QY 1 IHHIILECV 9
Db 1 IHHIILECV 9
```

```
RESULT 6
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from B6 and/or E7
; TITLE OF INVENTION: papillomavirus proteins and uses thereof
; FILE REFERENCE: 45636-5071-US
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide B6 15-44
US-10-476-570-53
```

```
Query Match          100.0%; Score 48; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHHIILECV 9
Db 16 IHHIILECV 24
```

```
RESULT 7
US-10-858-384-4
; Sequence 4, Application US/10858384
; Publication No. US20050030325A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
```

APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCINE
APPLICANT: FERRIS, ESTELLE
TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
TITLE OF INVENTION: PARTICULARLY IN VACCINATION
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858.384
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 4
LENGTH: 30
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4

Query Match 100.0%; Score 48; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHHIILBCV 9
Db 16 IHHIILBCV 24

RESULT 8
US-10-476-570-9
Sequence 9, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
TITLE OF INVENTION: Papillomavirus proteins and uses thereof
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 9
LENGTH: 32
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9

Query Match 100.0%; Score 48; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHHIILBCV 9
Db 17 IHHIILBCV 25

RESULT 9
US-10-476-570-19

Sequence 19, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLIERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
TITLE OF INVENTION: Papillomavirus proteins and uses thereof
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
CURRENT FILING DATE: 2003-11-04
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 19
LENGTH: 33
TYPE: PRT
ORGANISM: artificial sequence
FEATURE:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19

Query Match 100.0%; Score 48; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHHIILBCV 9
Db 17 IHHIILBCV 25

RESULT 10
US-10-177-390-6
Sequence 6, Application US/10177390
Publication No. US20030143743A1
GENERAL INFORMATION:
APPLICANT: Schuler, Gerold
APPLICANT: N.V. Antwerpse Innovatiecentrum
TITLE OF INVENTION: Improved Transfection of Eucaryotic Cells with Linear
TITLE OF INVENTION: Polynucleotides by Electroporation
FILE REFERENCE: 021505wo/JH/ml
CURRENT APPLICATION NUMBER: US/10/177,390
CURRENT FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-177-390-6

Query Match 100.0%; Score 48; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHHIILBCV 9
Db 23 IHHIILBCV 31

RESULT 11
US-10-484-063-20
Sequence 20, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILBERMO

```
APPLICANT: POLLEN, MICHELE
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: US-08-1037-1
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US/10/858,384
PRIOR FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2
```

```
Query Match      100.0%; Score 48; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
Qy      1 IHDIILECV 9
Db      23 IHDIILECV 31
```

```
RESULT 12
US-10-484-063-27
```

```
Sequence 27, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLERMO
APPLICANT: POLLEN, MICHELE
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: US-08-1037-1
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: US/10/484,063
PRIOR FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-484-063-27
```

```
Query Match      100.0%; Score 48; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.56;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      1 IHDIILECV 9
Db      23 IHDIILECV 31
```

```
RESULT 13
US-10-858-384-2
```

```
Sequence 2, Application US/10858384
Publication No. US20050033025A1
GENERAL INFORMATION:
APPLICANT: CHOPPIN, JEANNINE
APPLICANT: BOURGAULT VILLADA, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCES
APPLICANT: FERRIES, ESTELLE
TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
```

```
TITLE OF INVENTION: PARTICULARLY IN VACCINATION
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858,384
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2
```

```
Query Match      100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.59;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      1 IHDIILECV 9
Db      30 IHDIILECV 38
```

```
RESULT 14
US-10-367-057-16
```

```
Sequence 16, Application US/10367057
Publication No. US20050100554A1
GENERAL INFORMATION:
APPLICANT: Cuthill, Scott;
APPLICANT: Jackson, Amanda;
APPLICANT: Lewin, David A.;
APPLICANT: Ooi, Chean Eng
TITLE OF INVENTION: Complexes and Methods of Using Same
FILE REFERENCE: 21402-559
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: 60/256,911
PRIOR FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 198
SOFTWARE: Curoseqqlst version 0.1
SEQ ID NO 16
LENGTH: 158
TYPE: PRT
ORGANISM: Homo sapiens
US-10-367-057-16
```

```
Query Match      100.0%; Score 48; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.59;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      1 IHDIILECV 9
Db      30 IHDIILECV 38
```

```
RESULT 15
US-11-021-949-13
```

```
Sequence 13, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
PRIOR FILING DATE: 2003-12-23
NUMBER OF SEQ ID NOS: 361
```

```
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13
```

```
Query Match          100.0%; Score 48; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.59;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHDIILECV 9
Db 30 IHDIILECV 38
```

```
RESULT 16
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; PRIOR FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03271
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Construct
US-10-472-724-2
```

```
Query Match          100.0%; Score 48; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 0.64;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHDIILECV 9
Db 35 IHDIILECV 43
```

```
RESULT 17
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENV, Marie-Paule
; APPLICANT: BALLOU, Jean-Marc
; APPLICANT: BIZOUANE, Nadine
; TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; PRIOR FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
; SEQ ID NO 1
; LENGTH: 243
```

```
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone E6*TMF.
US-11-072-288-1
```

```
Query Match          100.0%; Score 48; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.92;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHDIILECV 9
Db 58 IHDIILECV 66
```

```
RESULT 18
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Publication No. US20020081329A1
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALLIAROS, JIM
; TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; PRIOR FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1
```

```
Query Match          100.0%; Score 48; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 IHDIILECV 9
Db 30 IHDIILECV 38
```

```
RESULT 19
US-10-000-903-4
; Sequence 4, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Gislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; PRIOR FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Homo sapien
```

US-10-000-903-4

Query Match 100.0%; Score 48; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
| | | | |
| | | | |
Db 136 IHDIILECV 144

RESULT 20

US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuncted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: Influenzae B and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4

Query Match 100.0%; Score 48; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
| | | | |
| | | | |
Db 136 IHDIILECV 144

RESULT 21

US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernandez
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT

; ORGANISM: Homo sapien
US-10-000-903-10

Query Match 100.0%; Score 48; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
| | | | |
| | | | |
Db 155 IHDIILECV 163

RESULT 22

US-10-899-771-10
; Sequence 10, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuncted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-10

Query Match 100.0%; Score 48; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
| | | | |
| | | | |
Db 155 IHDIILECV 163

RESULT 23

US-10-000-903-6
; Sequence 6, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernandez
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Benchelkh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371

TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-6

Query Match
Best Local Similarity 100.0%; Score 48; DB 4; Length 371;
Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
|||||
Db 136 IHDIILECV 144

RESULT 24
US-10-899-771-6
Sequence 6, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimaeic protein (protein D from Haemophilus
OTHER INFORMATION: Influenzae B and B6E7 fusion from Human Papilloma
OTHER INFORMATION: virus type 16)
US-10-899-771-6

Query Match
Best Local Similarity 100.0%; Score 48; DB 5; Length 371;
Pred. No. 1.4;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
|||||
Db 136 IHDIILECV 144

RESULT 25
US-10-000-903-14
Sequence 14, Application US/10000903
Publication No. US20020182221A1
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/10/000,903
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14

LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-10-000-903-14

Query Match
Best Local Similarity 100.0%; Score 48; DB 4; Length 390;
Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
|||||
Db 155 IHDIILECV 163

RESULT 26
US-10-899-771-14
Sequence 14, Application US/10899771
Publication No. US20050031638A1
GENERAL INFORMATION:
APPLICANT: Dalemans, Wilfried L.J.
APPLICANT: Gerard, Catherine Marie Ghislaine
TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
FILE REFERENCE: B45124
CURRENT APPLICATION NUMBER: US/10/899,771
CURRENT FILING DATE: 2004-07-27
PRIOR APPLICATION NUMBER: US/09/581,976
PRIOR FILING DATE: 2000-06-20
PRIOR APPLICATION NUMBER: PCT/EP98/08563
PRIOR FILING DATE: 1998-12-18
PRIOR APPLICATION NUMBER: GB 9727262.9
PRIOR FILING DATE: 1997-12-24
NUMBER OF SEQ ID NOS: 28
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Chimaeic protein (Clyta from Streptococcus
OTHER INFORMATION: pneumoniae and B6E7 fusion from Human papilloma
OTHER INFORMATION: virus type 16)
US-10-899-771-14

Query Match
Best Local Similarity 100.0%; Score 48; DB 5; Length 390;
Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
|||||
Db 155 IHDIILECV 163

RESULT 27
US-10-367-095-10
Sequence 10, Application US/10367095
Publication No. US20030228696A1
GENERAL INFORMATION:
APPLICANT: Robin A. Robinson
TITLE OF INVENTION: No. US20030228696A1 Insect Cell Line
FILE REFERENCE: 44149-1US1
CURRENT APPLICATION NUMBER: US/10/367,095
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,161
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,118
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,133
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,157
PRIOR FILING DATE: 2002-02-14

PRIOR APPLICATION NUMBER: US 60/356,156
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,123
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,113
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,154
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,135
PRIOR FILING DATE: 2002-02-14
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 10
LENGTH: 536
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-367-095-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
|||||
Db 500 IHDIILECV 508

RESULT 28
US-10-368-046-10

Sequence 10, Application US/10368046
Publication No. US20040063188A1
GENERAL INFORMATION:
APPLICANT: Robin A. Robinson
TITLE OF INVENTION: Method for Isolation and Purification of
FILE REFERENCE: 44149-3US1
CURRENT APPLICATION NUMBER: US/10/368,046
CURRENT FILING DATE: 2003-02-15
PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,161
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,118
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,133
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,157
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,156
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,123
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,113
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,154
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,135
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 10
LENGTH: 536
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-368-046-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
|||||
Db 500 IHDIILECV 508

RESULT 29
US-10-367-367-10

Sequence 10, Application US/10367367
Publication No. US20040121465A1
GENERAL INFORMATION:
APPLICANT: Robin A. Robinson
TITLE OF INVENTION: Optimization of Gene Sequences of
FILE REFERENCE: 44149-2US1
CURRENT APPLICATION NUMBER: US/10/367,367
CURRENT FILING DATE: 2003-02-15
PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,161
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,118
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,133
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,157
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,156
PRIOR FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 13
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 10
LENGTH: 536
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: HPV-16 L2/B6 fusion protein
US-10-367-367-10

Query Match 100.0%; Score 48; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIILECV 9
|||||
Db 500 IHDIILECV 508

RESULT 30
US-10-918-337-10

Sequence 10, Application US/10918337
Publication No. US20050118191A1
GENERAL INFORMATION:
APPLICANT: NOVAVAX, INC., et al.
TITLE OF INVENTION: Optimization of Gene Sequences of
FILE REFERENCE: 19065/2132
CURRENT APPLICATION NUMBER: US/10/918,337
CURRENT FILING DATE: 2004-08-13
PRIOR APPLICATION NUMBER: PCT/US03/04473
PRIOR FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: US 60/356,119
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,161
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,118
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,133
PRIOR FILING DATE: 2002-02-14
PRIOR APPLICATION NUMBER: US 60/356,157

```
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,156
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,123
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,113
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; PRIOR FILING DATE: 2002-02-14
;; PRIOR APPLICATION NUMBER: US 60/356,154
;; Remaining Prior Application data removed - See file wrapper or PALM.
;; NUMBER OF SEQ ID NOS: 13
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 10
;; LENGTH: 536
;; TYPE: PRT
;; ORGANISM: Artificial Sequence
;; FEATURE:
;; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10
```

```
Query Match          100.0%; Score 48; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 2.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 IHDIILBCV 9
Db 500 IHDIILBCV 508
```

```
RESULT 31
US-10-777-053-547
; Sequence 547, Application US/10777053
; Publication No. US20040132088A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Diamond, David C.
; APPLICANT: Qiu, Zhiyong
; APPLICANT: Lei, Xiang-Dong
; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
; TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
; FILE REFERENCE: MANNK.022C1
; CURRENT APPLICATION NUMBER: US/10/777,053
; PRIOR FILING DATE: 2004-02-10
; PRIOR APPLICATION NUMBER: 10/292,413
; PRIOR FILING DATE: 2002-11-07
; PRIOR APPLICATION NUMBER: 60/336,968
; PRIOR FILING DATE: 2001-11-07
; NUMBER OF SEQ ID NOS: 979
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 547
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human Papillomavirus 16
US-10-777-053-547
```

```
Query Match          91.7%; Score 44; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 IHDIILBC 8
Db 2 IHDIILBC 9
```

```
RESULT 32
US-10-837-217-547
; Sequence 547, Application US/10837217
; Publication No. US20040203051A1
; GENERAL INFORMATION:
; APPLICANT: Simard, John J. L.
; APPLICANT: Diamond, David C.
; APPLICANT: Qiu, Zhiyong
; APPLICANT: Lei, Xiang-Dong
```

```
;; TITLE OF INVENTION: EXPRESSION VECTORS ENCODING EPITOPES OF
;; TARGET-ASSOCIATED ANTIGENS AND METHODS FOR THEIR DESIGN
;; FILE REFERENCE: MANNK.022C2
;; CURRENT APPLICATION NUMBER: US/10/837,217
;; PRIOR FILING DATE: 2004-04-30
;; PRIOR APPLICATION NUMBER: 10/292,413
;; PRIOR FILING DATE: 2002-11-07
;; PRIOR APPLICATION NUMBER: 60/336,968
;; PRIOR FILING DATE: 2001-11-07
;; NUMBER OF SEQ ID NOS: 979
;; SOFTWARE: FastSeq for Windows Version 4.0
;; SEQ ID NO 547
;; LENGTH: 9
;; TYPE: PRT
;; ORGANISM: Human Papillomavirus 16
US-10-837-217-547
```

```
Query Match          91.7%; Score 44; DB 4; Length 9;
Best Local Similarity 100.0%; Pred. No. 1.7e+06;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 IHDIILBC 8
Db 2 IHDIILBC 9
```

```
RESULT 33
US-10-476-570-24
; Sequence 24, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 24
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 31-45
US-10-476-570-24
```

```
Query Match          91.7%; Score 44; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 0.28;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 2 HDIILBCV 9
Db 1 HDIILBCV 8
```

```
RESULT 34
US-10-424-599-275411
; Sequence 275411, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
```

```

; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 275411
; LENGTH: 297
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_90717C.1.pcp
US-10-424-599-275411

Query Match      81.2%; Score 39; DB 4; Length 297;
Best Local Similarity 62.5%; Pred. No. 52;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy      1 IHDIILEC 8
        |||:|:|
Db      179 IHDLVIEC 186

RESULT 35
US-10-424-599-168507
; Sequence 168507, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 168507
; LENGTH: 298
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(298)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_123177C.1.pcp
US-10-424-599-168507

Query Match      81.2%; Score 39; DB 4; Length 298;
Best Local Similarity 62.5%; Pred. No. 53;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

Qy      1 IHDIILEC 8
        |||:|:|
Db      178 IHDLVIEC 185

RESULT 36
US-11-021-949-19
; Sequence 19, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
```

```

; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 19
; LENGTH: 148
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-19

Query Match      79.2%; Score 38; DB 6; Length 148;
Best Local Similarity 77.8%; Pred. No. 39;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      1 IHDIILECV 9
        |||:|:|
Db      24 IHDLINDCV 32

RESULT 37
US-10-425-115-193535
; Sequence 193535, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 193535
; LENGTH: 428
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(428)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_108088C.1.pcp
US-10-425-115-193535

Query Match      79.2%; Score 38; DB 4; Length 428;
Best Local Similarity 75.0%; Pred. No. 1.2e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy      2 HDIILECV 9
        |||:|:|
Db      26 HDIILRCI 33

RESULT 38
US-10-739-930-10846
; Sequence 10846, Application US/10739930
; Publication No. US20040216190A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH
; TITLE OF INVENTION: PLANTS AND USES THEREOF FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53377)B
; CURRENT APPLICATION NUMBER: US/10/739,930
; CURRENT FILING DATE: 2003-12-18
; NUMBER OF SEQ ID NOS: 11088
; SEQ ID NO 10846
; LENGTH: 554
; TYPE: PRT
; ORGANISM: Triticum aestivum
```

FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(554)
OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
OTHER INFORMATION: Clone ID: TRIAB-23APR03-C7327_1.p
US-10-739-930-10846

Query Match 77.1%; Score 37; DB 5; Length 554;
Best Local Similarity 62.5%; Pred. No. 2.3e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

OY 1 HDIILECV 8
DB 458 VHDVLEEC 465

RESULT 39
US-10-437-963-195430
Sequence 195430, Application US/10437963
Publication No. US20040123343A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
APPLICANT: Wu, Wei
APPLICANT: Boukharov, Andrey A.
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53221)B
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 195430
LENGTH: 601
TYPE: PRT
ORGANISM: Oryza sativa
FEATURE:
NAME/KEY: unsure
LOCATION: (1)..(601)
OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT4530_9137C.1.p
US-10-437-963-195430

Query Match 77.1%; Score 37; DB 4; Length 601;
Best Local Similarity 55.6%; Pred. No. 2.6e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 HDIILECV 9
DB 557 VHDVLEEC 565

RESULT 40
US-10-424-599-219049
Sequence 219049, Application US/10424599
Publication No. US20040031072A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
FILE REFERENCE: 38-21(53223)B
CURRENT APPLICATION NUMBER: US/10/424,599
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 285684
SEQ ID NO 219049

LENGTH: 58
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT3847_39829C.1.p
US-10-424-599-219049

Query Match 75.0%; Score 36; DB 4; Length 58;
Best Local Similarity 55.6%; Pred. No. 35;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 HDIILECV 9
DB 38 HDVLEEC 46

RESULT 41
US-10-029-386-33593
Sequence 33593, Application US/10029386
Publication No. US20030194704A1
GENERAL INFORMATION:
APPLICANT: Penn, Sharon G.
APPLICANT: Rank, David R.
APPLICANT: Hanzel, David K.
TITLE OF INVENTION: HUMAN GENOME-DERIVED SINGLE EXON NUCLEIC ACID PROBES USEFUL FOR GI
TITLE OF INVENTION: EXPRESSION ANALYSIS TWO
FILE REFERENCE: ABOMICA-X-2
CURRENT APPLICATION NUMBER: US/10/029,386
CURRENT FILING DATE: 2001-12-20
NUMBER OF SEQ ID NOS: 34288
SOFTWARE: Annomax Sequence Listing Engine vers. 1.1
SEQ ID NO 33593
LENGTH: 70
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
OTHER INFORMATION: MAP TO CHR9.1
OTHER INFORMATION: EXPRESSED IN PETAL LIVER, SIGNAL = 1.1
OTHER INFORMATION: EXPRESSED IN HEART, SIGNAL = 0.59
OTHER INFORMATION: EXPRESSED IN ADULT LIVER, SIGNAL = 0.79
OTHER INFORMATION: SWISSPROT HIT: Q92574, VALUE 2.00e-30
US-10-029-386-33593

Query Match 75.0%; Score 36; DB 4; Length 70;
Best Local Similarity 57.1%; Pred. No. 42;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

OY 2 HDIILECV 8
DB 16 HDVLEEC 22

RESULT 42
US-10-425-115-205547
Sequence 205547, Application US/10425115
Publication No. US20040214272A1
GENERAL INFORMATION:
APPLICANT: La Rosa, Thomas J.
APPLICANT: Kovalic, David K.
APPLICANT: Zhou, Yihua
APPLICANT: Cao, Yongwei
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
TITLE OF INVENTION: Plants
FILE REFERENCE: 38-21(53222)B
CURRENT APPLICATION NUMBER: US/10/425,115
CURRENT FILING DATE: 2003-04-28
NUMBER OF SEQ ID NOS: 369326
SEQ ID NO 205547
LENGTH: 83
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
NAME/KEY: unsure

```
LOCATION: (1)..(83)
OTHER INFORMATION: unsure at all Xaa locations
FEATURE:
OTHER INFORMATION: Clone ID: MRT4577_119045C.1.pep
US-10-425-115-205547
```

```
Query Match          75.0%; Score 36; DB 4; Length 83;
Best Local Similarity 75.0%; Pred. No. 50;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy 1 IHDTILEC 8
Db 53 IHGVILEC 60
```

```
RESULT 43
US-11-021-949-361
Sequence 361, Application US/11021949
Publication No. US20050142541A1
```

```
GENERAL INFORMATION:
APPLICANT: LU, PETER
```

```
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARS, MICHAEL P.
```

```
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
```

```
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
```

```
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
```

```
PRIOR APPLICATION NUMBER: 60/532,373
NUMBER OF SEQ ID NOS: 361
```

```
SOFTWARE: FastSeq for Windows Version 4.0
SEQ ID NO 361
```

```
LENGTH: 158
TYPE: PRT
ORGANISM: human papilloma virus (HPV)
```

```
US-11-021-949-361
```

```
Query Match          75.0%; Score 36; DB 6; Length 158;
Best Local Similarity 55.6%; Pred. No. 98;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy 1 IHDTILEC 9
Db 25 LMDITDCV 33
```

```
RESULT 44
US-10-425-114-58813
Sequence 58813, Application US/10425114
```

```
Publication No. US20040034888A1
GENERAL INFORMATION:
```

```
APPLICANT: Liu, Jindong
APPLICANT: Zhou, Yihua
```

```
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E.
```

```
APPLICANT: Tabaska, Jack E.
APPLICANT: Cao, Yongwei
```

```
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(5313)B
```

```
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
```

```
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 58813
LENGTH: 274
TYPE: PRT
ORGANISM: Zea mays
```

```
FEATURE:
OTHER INFORMATION: Clone ID: 700085966_FLI.pep
US-10-425-114-58813
```

```
Query Match          75.0%; Score 36; DB 4; Length 274;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy 1 IHDTILEC 8
Db 114 VHDIFLDC 121
```

```
RESULT 45
US-10-425-114-65774
Sequence 65774, Application US/10425114
```

```
Publication No. US20040034888A1
GENERAL INFORMATION:
```

```
APPLICANT: Liu, Jindong
APPLICANT: Zhou, Yihua
```

```
APPLICANT: Kovalic, David K.
APPLICANT: Screen, Steven E.
```

```
APPLICANT: Tabaska, Jack E.
APPLICANT: Cao, Yongwei
```

```
TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(5313)B
```

```
CURRENT APPLICATION NUMBER: US/10/425,114
CURRENT FILING DATE: 2003-04-28
```

```
NUMBER OF SEQ ID NOS: 73128
SEQ ID NO 65774
```

```
LENGTH: 274
TYPE: PRT
ORGANISM: Zea mays
```

```
FEATURE:
OTHER INFORMATION: Clone ID: 700464936_FLI.pep
```

```
US-10-425-114-65774
```

```
Query Match          75.0%; Score 36; DB 4; Length 274;
Best Local Similarity 62.5%; Pred. No. 1.7e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy 1 IHDTILEC 8
Db 114 VHDIFLDC 121
```

```
RESULT 46
US-10-437-963-145297
Sequence 145297, Application US/10437963
```

```
Publication No. US20040123343A1
GENERAL INFORMATION:
```

```
APPLICANT: La Rosa, Thomas J.
APPLICANT: Zhou, Yihua
```

```
APPLICANT: Kovalic, David K.
APPLICANT: Cao, Yongwei
```

```
APPLICANT: Wu, Wei
APPLICANT: Boulharov, Andrey A.
```

```
APPLICANT: Barbazuk, Brad
APPLICANT: Li, Ping
```

```
TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
FILE REFERENCE: 38-21(53221)B
```

```
CURRENT APPLICATION NUMBER: US/10/437,963
CURRENT FILING DATE: 2003-05-14
```

```
NUMBER OF SEQ ID NOS: 204966
SEQ ID NO 145297
LENGTH: 291
TYPE: PRT
ORGANISM: Oryza sativa
```

```
FEATURE:
OTHER INFORMATION: Clone ID: PAT_MRT4530_4602C.1.pep
US-10-437-963-145297
```

```
Query Match          75.0%; Score 36; DB 4; Length 291;
Best Local Similarity 57.1%; Pred. No. 1.8e+02;
```

Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 HDIILEC 8
|:|:|:|
Db 183 HDVITEC 189

RESULT 47
US-10-424-599-187372
; Sequence 187372, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 187372
; LENGTH: 294
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_140209C.1.pep
US-10-424-599-187372

Query Match 75.0%; Score 36; DB 4; Length 294;
Best Local Similarity 55.6%; Pred. No. 1.9e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILECV 9
|:|:|:|
Db 182 LHDIFLQCL 190

RESULT 48
US-10-425-114-52907
; Sequence 52907, Application US/10425114
; Publication No. US20040034888A1
; GENERAL INFORMATION:
; APPLICANT: Liu, Jingdong
; APPLICANT: Zhou, Yihua
; APPLICANT: Kovalic, David K.
; APPLICANT: Screen, Steven E
; APPLICANT: Tabaska, Jack E
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53313)B
; CURRENT APPLICATION NUMBER: US/10/425,114
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 73128
; SEQ ID NO 52907
; LENGTH: 294
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; OTHER INFORMATION: Clone ID: LIB3061-015-E4_FLI.pep
US-10-425-114-52907

Query Match 75.0%; Score 36; DB 4; Length 294;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
|:|:|:|
Db 134 VHDIFLDC 141

RESULT 49
US-10-767-701-36891
; Sequence 36891, Application US/10767701
; Publication No. US20040172684A1
; GENERAL INFORMATION:
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53535)B
; CURRENT APPLICATION NUMBER: US/10/767,701
; CURRENT FILING DATE: 2004-01-29
; NUMBER OF SEQ ID NOS: 63128
; SEQ ID NO 36891
; LENGTH: 297
; TYPE: PRT
; ORGANISM: Sorghum bicolor
; FEATURE:
; OTHER INFORMATION: Clone ID: SORBI-28MAY03-C120388_1.pep
US-10-767-701-36891

Query Match 75.0%; Score 36; DB 4; Length 297;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
|:|:|:|
Db 137 VHDIFLDC 144

RESULT 50
US-10-425-115-345578
; Sequence 345578, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 345578
; LENGTH: 304
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(304)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_78335C.1.pep
US-10-425-115-345578

Query Match 75.0%; Score 36; DB 4; Length 304;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
|:|:|:|
Db 192 VHDIAIEC 199

Search completed: May 5, 2006, 08:28:31
Udb time : 60 secs

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OM protein - protein search, using bw model

Run on: May 5, 2006, 08:19:00 ; Search time 8.4 Seconds
(without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-8
Perfect score: 48
Sequence: 1 IHDIILCECV 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

Published Applications AA_New:*
1: /SID5/prodata/1/pubppa/US08_NEW_PUB.pep1.*
2: /SID5/prodata/1/pubppa/US07_NEW_PUB.pep.*
3: /SID5/prodata/1/pubppa/US08_NEW_PUB.pep.*
4: /SID5/prodata/1/pubppa/US08_NEW_PUB.pep.*
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

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5	48	100.0	248	9	US-10-530-253-5
6	48	100.0	248	9	US-10-530-253-7
7	48	100.0	248	9	US-10-530-253-9
8	48	100.0	248	9	US-10-530-253-11
9	48	100.0	256	11	US-11-192-923A-2
10	38	79.2	10	9	US-10-530-061-53
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13	36	75.0	298	11	US-11-172-740-1141
14	36	75.0	306	11	US-11-172-740-1144
15	36	75.0	310	11	US-11-172-740-1142
16	35	72.9	158	9	US-10-530-253-26
17	35	72.9	174	11	US-11-096-568A-6616
18	35	72.9	220	11	US-11-087-099-4165
19	35	72.9	247	11	US-11-096-568A-6615
20	35	72.9	301	11	US-11-096-568A-6614
21	35	72.9	335	11	US-11-087-099-10727

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27	210	11	US-11-096-568A-24716	Sequence 24716, A
28	310	11	US-11-096-568A-21631	Sequence 21631, A
29	311	11	US-11-096-568A-21630	Sequence 21630, A
30	351	11	US-11-087-099-2242	Sequence 2242, Ap
31	351	11	US-11-087-099-8245	Sequence 8245, Ap
32	15	9	US-10-530-061-1673	Sequence 1673, Ap
33	148	9	US-10-530-253-22	Sequence 22, Appl
34	149	9	US-10-530-253-17	Sequence 17, Appl
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37	392	11	US-11-087-099-9859	Sequence 9859, Ap
38	623	11	US-11-188-298-16877	Sequence 16877, A
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40	149	9	US-10-530-253-24	Sequence 24, Appl
41	160	9	US-10-530-253-25	Sequence 25, Appl
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90	596	9	US-10-504-879-10	Sequence 10, Appl
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93	630	9	US-10-504-879-26	Sequence 26, Appl
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256	29	60.4	840	9	US-11-239-674-90	Sequence 90, App1	329	28	58.3	623	9	US-10-793-626-1068	Sequence 1068, Ap
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258	29	60.4	840	9	US-10-725-475-16	Sequence 16, App1	331	28	58.3	651	9	US-10-517-939-268	Sequence 268, App
259	29	60.4	842	9	US-10-645-441-2	Sequence 2, App11	332	28	58.3	655	11	US-11-124-368A-262	Sequence 262, App
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262	29	60.4	1125	11	US-11-024-959-360	Sequence 360, App	335	28	58.3	700	11	US-11-087-099-5110	Sequence 5110, Ap
263	29	60.4	1221	9	US-10-858-730-222	Sequence 222, App	336	28	58.3	738	9	US-10-995-561-692	Sequence 692, App
264	29	60.4	1263	9	US-10-330-773-539	Sequence 539, App1	337	28	58.3	738	9	US-10-995-561-693	Sequence 693, App
265	29	60.4	1463	11	US-11-080-991-22	Sequence 22, App1	338	28	58.3	738	9	US-10-880-881-23	Sequence 881, Ap
266	29	60.4	1619	11	US-11-087-099-10709	Sequence 10709, A	339	28	58.3	810	9	US-10-453-372-1116	Sequence 4218, Ap
267	29	60.4	2333	11	US-11-096-281-13	Sequence 13, App1	340	28	58.3	825	11	US-11-087-099-5521	Sequence 5521, Ap
268	29	60.4	2339	11	US-11-096-281-11	Sequence 11, App1	341	28	58.3	825	11	US-11-087-099-5521	Sequence 5521, Ap
269	28	58.3	66	9	US-10-467-657-6624	Sequence 6624, Ap	342	28	58.3	825	11	US-11-188-298-5094	Sequence 5094, Ap
270	28	58.3	83	8	US-10-501-834-21	Sequence 21, App1	343	28	58.3	825	11	US-11-188-298-13018	Sequence 13018, A
271	28	58.3	103	11	US-11-188-298-14282	Sequence 14282, A	344	28	58.3	855	11	US-11-087-099-10166	Sequence 20399, A
272	28	58.3	109	11	US-11-188-298-11984	Sequence 11984, A	345	28	58.3	855	11	US-11-188-298-20395	Sequence 20395, A
273	28	58.3	124	9	US-10-510-386-140	Sequence 140, App	346	28	58.3	855	11	US-11-144-987-16	Sequence 2532, Ap
274	28	58.3	124	9	US-10-467-657-2682	Sequence 2682, Ap	347	28	58.3	915	11	US-11-144-987-16	Sequence 16, App1
275	28	58.3	124	11	US-11-045-004-2827	Sequence 2827, Ap	348	28	58.3	915	11	US-11-144-987-16	Sequence 16, App1
276	28	58.3	138	11	US-11-188-298-19697	Sequence 19697, A	349	28	58.3	915	11	US-11-144-987-16	Sequence 16, App1
277	28	58.3	139	11	US-11-096-568A-16621	Sequence 16621, A	350	28	58.3	915	11	US-11-144-987-16	Sequence 16, App1
278	28	58.3	235	9	US-10-467-657-654	Sequence 654, App	351	28	58.3	915	11	US-11-144-987-16	Sequence 16, App1
279	28	58.3	241	11	US-11-188-298-22369	Sequence 22369, A	352	28	58.3	917	11	US-11-144-987-20	Sequence 20, App1
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282	28	58.3	287	11	US-11-079-463-6784	Sequence 6784, Ap	355	28	58.3	917	11	US-11-144-987-26	Sequence 26, App1
283	28	58.3	292	11	US-11-079-463-5410	Sequence 5410, Ap	356	28	58.3	917	11	US-11-205-935-18	Sequence 18, App1
284	28	58.3	296	11	US-11-188-298-21966	Sequence 21966, A	357	28	58.3	917	11	US-11-205-935-24	Sequence 24, App1
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287	28	58.3	320	11	US-11-087-099-7772	Sequence 7772, Ap	360	28	58.3	984	11	US-11-113-424-61	Sequence 85, App1
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289	28	58.3	322	11	US-11-188-298-6157	Sequence 6157, Ap	362	28	58.3	985	11	US-11-113-424-61	Sequence 85, App1
290	28	58.3	325	11	US-11-188-298-6157	Sequence 6157, Ap	363	28	58.3	986	11	US-11-203-551A-80	Sequence 80, App1
291	28	58.3	325	11	US-11-098-686-10272	Sequence 10272, A	364	28	58.3	987	9	US-10-949-720-395	Sequence 395, App
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296	28	58.3	345	11	US-11-188-298-12944	Sequence 12944, A	369	28	58.3	998	11	US-11-203-551A-89	Sequence 89, App1
297	28	58.3	347	11	US-11-087-099-11600	Sequence 11600, A	370	28	58.3	998	11	US-11-203-551A-89	Sequence 89, App1
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299	28	58.3	349	11	US-11-087-099-3545	Sequence 3545, Ap	372	28	58.3	1005	11	US-11-203-551A-84	Sequence 84, App1
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305	28	58.3	389	11	US-11-188-298-14197	Sequence 14197, A	378	28	58.3	1055	11	US-11-051-120-1417	Sequence 1017, Ap
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309	28	58.3	408	11	US-11-069-642-115	Sequence 115, App	382	28	58.3	1115	9	US-10-195-883-440	Sequence 440, App
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395	28	58.3	1441	11	US-11-096-568A-28129	Sequence 28129, A	468	27	56.2	308	11	US-11-096-568A-35812	Sequence 25812, A
396	28	58.3	1461	11	US-11-183-136-28	Sequence 28, Appl1	469	27	56.2	310	11	US-11-096-568A-30961	Sequence 30961, A
397	28	58.3	1490	11	US-11-096-568A-28128	Sequence 28128, A	470	27	56.2	311	11	US-11-188-298-13038	Sequence 13038, A
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402	28	58.3	3507	11	US-11-075-185-7	Sequence 7, Appl1	475	27	56.2	335	11	US-11-188-298-447	Sequence 447, App
403	27	57.3	509	9	US-10-506-454-1427	Sequence 1427, App	476	27	56.2	335	11	US-11-188-298-8546	Sequence 8546, App
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421	27	56.2	168	11	US-11-096-568A-32249	Sequence 32249, A	494	27	56.2	341	11	US-11-188-298-4610	Sequence 4610, App
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543	27	56.2	412	11	US-11-087-099-499	Sequence 499, App	616	26	54.2	14	11	US-11-004-399-3210	Sequence 3210, Ap
544	27	56.2	413	11	US-11-188-298-3235	Sequence 3235, Ap	617	26	54.2	40	11	US-11-129-741-3553	Sequence 3553, Ap
545	27	56.2	413	11	US-11-188-298-15320	Sequence 15320, A	618	26	54.2	70	11	US-11-096-568A-13282	Sequence 3282, Ap
546	27	56.2	414	11	US-11-174-341-160	Sequence 160, App	619	26	54.2	70	11	US-11-096-568A-3284	Sequence 3284, Ap
547	27	56.2	415	11	US-11-168-298-11955	Sequence 11955, A	620	26	54.2	80	9	US-10-467-657-9966	Sequence 5966, Ap
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549	27	56.2	417	11	US-11-188-298-2525	Sequence 2525, Ap	622	26	54.2	85	11	US-11-338-285A-2	Sequence 4536, Ap
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830	26	54.2	463	11	US-11-156-084-41	Sequence 41, App	903	26	54.2	704	9	US-10-505-263-10	Sequence 10, App
831	26	54.2	463	11	US-11-156-084-242	Sequence 242, App	904	26	54.2	718	11	US-11-188-298-14144	Sequence 14144, A
832	26	54.2	466	11	US-11-156-084-42	Sequence 42, App	905	26	54.2	718	11	US-11-188-298-18845	Sequence 18845, A
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838	26	54.2	475	11	US-11-096-568A-9140	Sequence 9140, App	911	26	54.2	757	11	US-11-031-206-184	Sequence 184, App
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842	26	54.2	483	11	US-11-188-298-17721	Sequence 17721, A	915	26	54.2	777	9	US-10-645-441-3	Sequence 441, App
843	26	54.2	484	9	US-10-467-657-5472	Sequence 5472, App	916	26	54.2	788	9	US-10-505-263-85	Sequence 85, App
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846	26	54.2	489	9	US-10-195-883-116	Sequence 116, App	919	26	54.2	841	11	US-11-050-804-2	Sequence 2, App
847	26	54.2	489	9	US-10-195-883-116	Sequence 116, App	920	26	54.2	861	11	US-11-087-099-1595	Sequence 1595, App
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851	26	54.2	492	9	US-10-216-161A-7	Sequence 7, App	924	26	54.2	886	9	US-10-821-234-1390	Sequence 1390, App
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866	26	54.2	511	11	US-11-096-568A-3307	Sequence 3307, App	939	26	54.2	1036	9	US-10-243-116-104	Sequence 104, App
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772 26 54.2 1706 11 US-11-086-568A-27854 Sequence 27854, A
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774 26 54.2 1767 9 US-10-204-639-54 Sequence 54, Appl
775 26 54.2 1896 9 US-10-877-346-13 Sequence 13, Appl
776 26 54.2 1963 9 US-10-877-346-13 Sequence 43, Appl
777 26 54.2 2004 9 US-10-469-469-250 Sequence 250, App
778 26 54.2 2215 8 US-10-505-928-310 Sequence 310, App
779 26 54.2 2376 11 US-11-096-568A-27513 Sequence 27513, A
780 26 54.2 2410 11 US-11-175-689-8 Sequence 8, Appl
781 26 54.2 2518 11 US-11-096-568A-27512 Sequence 27512, A
782 26 54.2 2531 11 US-11-124-367A-467 Sequence 467, App
783 26 54.2 2553 11 US-11-096-568A-27511 Sequence 27511, A
784 26 54.2 3012 11 US-11-124-367A-465 Sequence 465, App
785 26 54.2 3144 11 US-11-055-035-1 Sequence 1, Appl
786 26 54.2 4097 9 US-10-501-035-263 Sequence 263, Appl
787 26 54.2 4128 9 US-10-770-726-77 Sequence 77, Appl
788 26 54.2 4443 11 US-11-129-741-3478 Sequence 3478, App
789 26 54.2 4473 9 US-10-895-064-460 Sequence 460, App
790 26 54.2 4473 11 US-11-129-741-460 Sequence 460, App
791 26 54.2 16990 11 US-11-175-689-7 Sequence 7, Appl
792 25.5 53.1 212 11 US-11-098-686-11253 Sequence 11253, A
793 25.5 53.1 338 11 US-11-087-099-4569 Sequence 4569, App
794 25.5 53.1 783 11 US-11-188-298-9347 Sequence 9347, App
795 25.5 53.1 826 10 US-11-301-554-330 Sequence 330, App
796 25.5 53.1 826 10 US-11-288-720-10 Sequence 10, Appl
797 25.5 53.1 1047 11 US-11-124-367A-388 Sequence 388, App
798 25.5 53.1 1058 11 US-11-124-367A-386 Sequence 386, App
799 25.5 53.1 1062 11 US-11-124-367A-387 Sequence 387, App
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ALIGNMENTS

RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13

Query Match 100.0%; Score 48; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.028; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 IHDIILCEV 9
Db 23 IHDIILCEV 31

RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1

; GENERAL INFORMATION:
; APPLICANT: Healthbanc Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; CURRENT FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3

Query Match 100.0%; Score 48; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.029; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 IHDIILCEV 9
Db 30 IHDIILCEV 38

RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-1

Query Match 100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.046; Indels 0; Gaps 0;
Matches 9; Conservative 0; Mismatches 0;

Qy 1 IHDIILCEV 9
Db 23 IHDIILCEV 31

RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929

;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7
Query Match
Best Local Similarity 100.0%; Score 48; DB 9; Length 248;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDITLCV 9
Db 23 IHDITLCV 31

RESULT 5
US-10-530-253-5
; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5

Query Match 100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.046;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDITLCV 9
Db 23 IHDITLCV 31

RESULT 6
US-10-530-253-7
; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 248

;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 16
US-10-530-253-7

Query Match
Best Local Similarity 100.0%; Score 48; DB 9; Length 248;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDITLCV 9
Db 120 IHDITLCV 128

RESULT 7
US-10-530-253-9
; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9

Query Match 100.0%; Score 48; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.046;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDITLCV 9
Db 120 IHDITLCV 128

RESULT 8
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11

Query Match 100.0%; Score 48; DB 9; Length 248;

Best Local Similarity 100.0%; Pred. No. 0.046;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIIIECV 9
Db 120 IHDIIIECV 128

RESULT 9

US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060019928A1
; GENERAL INFORMATION:
; APPLICANT: PANG, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; TITLE OF INVENTION: RECOMBINANT REPLICON
; FILE REFERENCE: 116620-003
; CURRENT FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

Query Match 100.0%; Score 48; DB 11; Length 256;
Best Local Similarity 100.0%; Pred. No. 0.048;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Qy 1 IHDIIIECV 9
Db 128 IHDIIIECV 136

RESULT 10

US-10-530-061-53
; Sequence 53, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 53
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-53

Query Match 79.2%; Score 38; DB 9; Length 10;
Best Local Similarity 88.9%; Pred. No. 0.16;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 IHDIIIECV 9
Db 1 IHDIIIECV 9

RESULT 11
US-10-530-061-114
; Sequence 114, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 114
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-114

Query Match 79.2%; Score 38; DB 9; Length 10;
Best Local Similarity 88.9%; Pred. No. 0.16;
Matches 8; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 1 IHDIIIECV 9
Db 1 IHDIIIECV 9

RESULT 12

US-11-172-740-1143
; Sequence 1143, Application US/11172740
; Publication No. US20060057724A1
; GENERAL INFORMATION:
; APPLICANT: MASCIA, Peter
; APPLICANT: ALEXANDROV, Nikolai
; APPLICANT: BROVER, Vyacheslav
; TITLE OF INVENTION: NOCLOTTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR ;
; FILE REFERENCE: 2750-1602PUS2
; CURRENT APPLICATION NUMBER: US/11/172,740
; CURRENT FILING DATE: 2005-06-30
; PRIOR APPLICATION NUMBER: 60/583,621
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,829
; PRIOR FILING DATE: 2004-06-30
; PRIOR APPLICATION NUMBER: 60/584,800
; PRIOR FILING DATE: 2004-06-30
; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1143
; LENGTH: 202
; TYPE: PRT
; ORGANISM: Zea mays

NAME/KEY: misc_feature
LOCATION: (1)..(202)
OTHER INFORMATION: Public GI no. 34903270
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Useful for increasing chlorophyll and photosynthetic cap
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Useful for making lethal plants for genetic confinement


```
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with increased biomass
US-11-172-740-1143
Query Match          75.0%; Score 36; DB 11; Length 202;
Best Local Similarity 62.5%; Pred. No. 8.6;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 IHD11LEC 8
Db 42 VHD1FLDC 49

RESULT 13
US-11-172-740-1141
Sequence 1141, Application US/11172740
Publication No. US20060057724A1
GENERAL INFORMATION:
APPLICANT: MASCTA, Peter
APPLICANT: ALEXANDROV, Nickolai
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
FILE REFERENCE: 2750-1602PUS2
CURRENT FILING DATE: 2005-06-30
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,621
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,829
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,800
PRIOR FILING DATE: 2004-06-30
NUMBER OF SEQ ID NOS: 2523
SEQ ID NO 1141
LENGTH: 298
TYPE: PRT
ORGANISM: Triticum aestivum
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1) .. (298)
OTHER INFORMATION: Ceres CLONE ID no. 788296
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
FEATURE:
NAME/KEY: misc_feature
```

```
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with increased biomass
US-11-172-740-1141
Query Match          75.0%; Score 36; DB 11; Length 298;
Best Local Similarity 62.5%; Pred. No. 13;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

Qy 1 IHD11LEC 8
Db 138 VHD1FLDC 145

RESULT 14
US-11-172-740-1144
Sequence 1144, Application US/11172740
Publication No. US20060057724A1
GENERAL INFORMATION:
APPLICANT: MASCTA, Peter
APPLICANT: ALEXANDROV, Nickolai
APPLICANT: BROVER, Vyacheslav
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
FILE REFERENCE: 2750-1602PUS2
CURRENT FILING DATE: 2005-06-30
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/583,621
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,829
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,800
PRIOR FILING DATE: 2004-06-30
NUMBER OF SEQ ID NOS: 2523
SEQ ID NO 1144
LENGTH: 306
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1) .. (306)
OTHER INFORMATION: Public GI no. 56783703
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with altered leaf shape eg curl
```

FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with increased biomass
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making smaller plants
US-11-172-740-1144

Query Match 75.0%; Score 36; DB 11; Length 306;
Best Local Similarity 62.5%; Pred. No. 13;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIILBC 8
Db 146 VHDIFLDC 153

RESULT 15
US-11-172-740-1142
Sequence 1142, Application US/11172740
Publication No. US2006005724A1
GENERAL INFORMATION:
APPLICANT: MASCIA, Peter
APPLICANT: ALEXANDROV, Nickolai
TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
FILE REFERENCE: 2750-1602PUS2
CURRENT APPLICATION NUMBER: US/11/172,740
PRIOR FILING DATE: 2005-06-30
PRIOR APPLICATION NUMBER: 60/583,621
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,829
PRIOR FILING DATE: 2004-06-30
PRIOR APPLICATION NUMBER: 60/584,800
PRIOR FILING DATE: 2004-06-30
NUMBER OF SEQ ID NOS: 2523
SEQ ID NO 1142
LENGTH: 310
TYPE: PRT
ORGANISM: Zea mays
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(310)
OTHER INFORMATION: Ceres CLONE ID no. 303545
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
FEATURE:
NAME/KEY: misc_feature
LOCATION:
OTHER INFORMATION: Utility: Useful for making plants with increased biomass
FEATURE:
NAME/KEY: misc_feature

LOCATION:
OTHER INFORMATION: Utility: Useful for making smaller plants
US-11-172-740-1142

Query Match 75.0%; Score 36; DB 11; Length 310;
Best Local Similarity 62.5%; Pred. No. 13;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIILBC 8
Db 150 VHDIFLDC 157

RESULT 16
US-10-530-253-26
Sequence 26, Application US/10530253
Publication No. US20060014926A1
GENERAL INFORMATION:
APPLICANT: Casaretti, Maria C.
APPLICANT: Smith, Larry
APPLICANT: Jeffrey K. Pullen
APPLICANT: Susan P. McElhinney
TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
FILE REFERENCE: 00630/100M137-US2
CURRENT APPLICATION NUMBER: US/10/530,253
CURRENT FILING DATE: 2005-04-04
PRIOR APPLICATION NUMBER: PCT/US2003/031726
PRIOR FILING DATE: 2003-10-02
PRIOR APPLICATION NUMBER: US 60/415,929
PRIOR FILING DATE: 2002-10-03
NUMBER OF SEQ ID NOS: 65
SOFTWARE: PatentIn version 3.1
SEQ ID NO 26
LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 68
US-10-530-253-26

Query Match 72.9%; Score 35; DB 9; Length 158;
Best Local Similarity 44.4%; Pred. No. 10;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIILBCV 9
Db 25 LHDVTIDCV 33

RESULT 17
US-11-096-568A-6616
Sequence 6616, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nickolai et al.
TITLE OF INVENTION: Sequence-determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 6616
LENGTH: 174
TYPE: PRT
ORGANISM: Glycine max
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)..(174)
OTHER INFORMATION: Ceres Seq. ID no. 14315992
US-11-096-568A-6616

Query Match 72.9%; Score 35; DB 11; Length 174;
Best Local Similarity 50.0%; Pred. No. 12;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIEC 8
: ||: ||
Db 54 VHDVITEC 61

RESULT 18

US-11-087-099-4165
; Sequence 4165, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 4165
; LENGTH: 220
; TYPE: PRT
; ORGANISM: Trifolium aestivum
US-11-087-099-4165

Query Match 72.9%; Score 35; DB 11; Length 220;
Best Local Similarity 57.1%; Pred. No. 15;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 HDIIEC 8
: ||: ||
Db 111 HDVITEC 117

RESULT 19

US-11-096-568A-6615
; Sequence 6615, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6615
; LENGTH: 247
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)-(247)
; OTHER INFORMATION: Ceres Seq. ID no. 14315991
US-11-096-568A-6615

Query Match 72.9%; Score 35; DB 11; Length 247;
Best Local Similarity 50.0%; Pred. No. 17;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIEC 8
: ||: ||
Db 127 VHDVITEC 134

RESULT 20

US-11-096-568A-6614
; Sequence 6614, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01

NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 6614
; LENGTH: 301
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)-(301)
; OTHER INFORMATION: Ceres Seq. ID no. 14315990
US-11-096-568A-6614

Query Match 72.9%; Score 35; DB 11; Length 301;
Best Local Similarity 50.0%; Pred. No. 20;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIEC 8
: ||: ||
Db 181 VHDVITEC 188

RESULT 21

US-11-087-099-10727
; Sequence 10727, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 10727
; LENGTH: 335
; TYPE: PRT
; ORGANISM: Thermoplasma volcanium
US-11-087-099-10727

Query Match 72.9%; Score 35; DB 11; Length 335;
Best Local Similarity 85.7%; Pred. No. 23;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 DIIECV 9
: ||: ||
Db 230 DIVIECV 236

RESULT 22

US-11-087-099-12277
; Sequence 12277, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 12277
; LENGTH: 336
; TYPE: PRT
; ORGANISM: Thermoplasma acidophilum
US-11-087-099-12277

Query Match 72.9%; Score 35; DB 11; Length 336;
Best Local Similarity 85.7%; Pred. No. 23;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 3 DIIECV 9
: ||: ||
Db 230 DIVIECV 236

RESULT 23

```
US-10-858-730-70
; Sequence 70, Application US/10858730
; Publication No. US20050255568A1
; GENERAL INFORMATION:
; APPLICANT: Bailey, Richard B.
; APPLICANT: Blomquist, Paul
; APPLICANT: Doten, Reed
; APPLICANT: Driggers, Edward M.
; APPLICANT: Madden, Kevin T.
; APPLICANT: O'Leary, Jessica
; APPLICANT: O'Toole, George
; APPLICANT: Trueheart, Joshua
; APPLICANT: Walbridge, Michael J.
; APPLICANT: Yorgey, Peter S.
; TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR AMINO ACID
; FILE REFERENCE: 14184-030001
; CURRENT APPLICATION NUMBER: US/10/858,730
; CURRENT FILING DATE: 2004-06-01
; PRIOR APPLICATION NUMBER: US 60/475,000
; PRIOR FILING DATE: 2003-05-30
; PRIOR APPLICATION NUMBER: US 60/551,860
; PRIOR FILING DATE: 2004-03-10
; NUMBER OF SEQ ID NOS: 364
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 70
; LENGTH: 1158
; TYPE: PRT
; ORGANISM: Thermobifida fusca
US-10-858-730-70
```

```
Query Match          72.9%; Score 35; DB 9; Length 1158;
Best Local Similarity 62.5%; Pred. No. 81;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      2 HDIILECV 9
Db      498 HDIIVDC 505
```

```
RESULT 24
US-10-530-253-18
; Sequence 18, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan F. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 35
US-10-530-253-18
```

```
Query Match          70.8%; Score 34; DB 9; Length 149;
Best Local Similarity 66.7%; Pred. No. 15;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;
```

```
Qy      1 HDIILCECV 9
Db      23 IHEICLNCV 31
```

```
RESULT 25
US-11-096-568A-24717
; Sequence 24717, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24717
; LENGTH: 153
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(153)
; OTHER INFORMATION: Cereals Seq. ID no. 12450486
US-11-096-568A-24717
```

```
Query Match          70.8%; Score 34; DB 11; Length 153;
Best Local Similarity 57.1%; Pred. No. 16;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy      2 HDIILBC 8
Db      126 HDIIVDC 132
```

```
RESULT 26
US-11-096-568A-21632
; Sequence 21632, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 21632
; LENGTH: 204
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(204)
; OTHER INFORMATION: Cereals Seq. ID no. 12405549
US-11-096-568A-21632
```

```
Query Match          70.8%; Score 34; DB 11; Length 204;
Best Local Similarity 62.5%; Pred. No. 21;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
```

```
Qy      1 HDIILBC 8
Db      104 LHDISIEC 111
```

```
RESULT 27
US-11-096-568A-24716
; Sequence 24716, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nickolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
```

CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 24716
LENGTH: 210
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(210)
OTHER INFORMATION: Ceres Seq. ID no. 12450485
US-11-096-568A-24716

Query Match
Best Local Similarity 57.1%; Score 34; DB 11; Length 210;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 2 HDIILEC 8
DB 183 HDIVIDC 189

RESULT 28
US-11-096-568A-21631
Sequence 21631, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 21631
LENGTH: 310
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(310)
OTHER INFORMATION: Ceres Seq. ID no. 1240548
US-11-096-568A-21631

Query Match
Best Local Similarity 62.5%; Score 34; DB 11; Length 310;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
DB 210 LHDISIEC 217

RESULT 29
US-11-096-568A-21630
Sequence 21630, Application US/11096568A
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 21630
LENGTH: 311
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)..(311)
OTHER INFORMATION: Ceres Seq. ID no. 12405547
US-11-096-568A-21630

Query Match
Best Local Similarity 70.8%; Score 34; DB 11; Length 311;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
DB 211 LHDISIEC 218

RESULT 30
US-11-087-099-2242
Sequence 2242, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B BP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 2242
LENGTH: 351
TYPE: PRT
ORGANISM: Streptococcus agalactiae NEM316
US-11-087-099-2242

Query Match
Best Local Similarity 71.4%; Score 34; DB 11; Length 351;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 3 DIILECV 9
DB 238 DVILECV 244

RESULT 31
US-11-087-099-8245
Sequence 8245, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B BP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 8245
LENGTH: 351
TYPE: PRT
ORGANISM: Streptococcus agalactiae 2603V/R
US-11-087-099-8245

Query Match
Best Local Similarity 71.4%; Score 34; DB 11; Length 351;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 3 DIILECV 9
DB 238 DVILECV 244

RESULT 32
US-10-530-061-1673
Sequence 1673, Application US/10530061
Publication No. US20060079453A1
GENERAL INFORMATION:
APPLICANT: SIDNEY, JOHN
APPLICANT: SOUTHWOOD, SCOTT
APPLICANT: SETTE, ALESSANDRO
TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
FILE REFERENCE: 2060.03US02/EKS/M-M
CURRENT APPLICATION NUMBER: US/10/530,061

;; CURRENT FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US03/31308
;; PRIOR FILING DATE: 2003-10-03
;; PRIOR APPLICATION NUMBER: 60/416,207
;; PRIOR FILING DATE: 2002-10-03
;; PRIOR APPLICATION NUMBER: 60/417,269
;; PRIOR FILING DATE: 2002-10-08
;; NUMBER OF SEQ ID NOS: 2503
;; SOFTWARE: PatentIn version 3.3
;; SEQ ID NO 1673
;; LENGTH: 15
;; TYPE: PRT
;; ORGANISM: Human papillomavirus
US-10-530-061-1673

Query Match 68.8%; Score 33; DB 9; Length 15;
Best Local Similarity 66.7%; Pred. No. 2.3;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIECV 9
||:|:|
Db 4 IHNIELQCV 12

RESULT 33
US-10-530-253-22
;; Sequence 22, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Casasetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; APPLICANT: Susan P. McElhinney
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M17-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; CURRENT FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02
;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 22
;; LENGTH: 148
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 52
US-10-530-253-22

Query Match 68.8%; Score 33; DB 9; Length 148;
Best Local Similarity 55.6%; Pred. No. 24;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIECV 9
||:|:|
Db 23 VHEIRLQCV 31

RESULT 34
US-10-530-253-17
;; Sequence 17, Application US/10530253
;; Publication No. US20060014926A1
;; GENERAL INFORMATION:
;; APPLICANT: Casasetti, Maria C.
;; APPLICANT: Smith, Larry
;; APPLICANT: Jeffrey K. Pullen
;; APPLICANT: Susan P. McElhinney
;; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
;; FILE REFERENCE: 00630/100M17-US2
;; CURRENT APPLICATION NUMBER: US/10/530,253
;; CURRENT FILING DATE: 2005-04-04
;; PRIOR APPLICATION NUMBER: PCT/US2003/031726
;; PRIOR FILING DATE: 2003-10-02

;; PRIOR APPLICATION NUMBER: US 60/415,929
;; PRIOR FILING DATE: 2002-10-03
;; NUMBER OF SEQ ID NOS: 65
;; SOFTWARE: PatentIn version 3.1
;; SEQ ID NO 17
;; LENGTH: 149
;; TYPE: PRT
;; ORGANISM: Human papillomavirus type 33
US-10-530-253-17

Query Match 68.8%; Score 33; DB 9; Length 149;
Best Local Similarity 66.7%; Pred. No. 24;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 IHDIIECV 9
||:|:|
Db 23 IHNIELQCV 31

RESULT 35
US-11-264-096-110
;; Sequence 110, Application US/11264096
;; Publication No. US20060084794A1
;; GENERAL INFORMATION:
;; APPLICANT: Rosen et al.
;; TITLE OF INVENTION: Albumin Fusion Proteins
;; FILE REFERENCE: P546D1
;; CURRENT APPLICATION NUMBER: US/11/264,096
;; CURRENT FILING DATE: 2005-11-02
;; PRIOR APPLICATION NUMBER: 09/833,245
;; PRIOR FILING DATE: 2001-04-12
;; PRIOR APPLICATION NUMBER: 60/229,358
;; PRIOR FILING DATE: 2000-04-12
;; PRIOR APPLICATION NUMBER: 60/256,931
;; PRIOR FILING DATE: 2000-12-21
;; PRIOR APPLICATION NUMBER: 60/199,384
;; PRIOR FILING DATE: 2000-04-25
;; NUMBER OF SEQ ID NOS: 2267
;; SOFTWARE: PatentIn Ver. 2.1
;; SEQ ID NO 110
;; LENGTH: 157
;; TYPE: PRT
;; ORGANISM: Homo sapiens
US-11-264-096-110

Query Match 68.8%; Score 33; DB 11; Length 157;
Best Local Similarity 71.4%; Pred. No. 26;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 1 IHDIIECV 7
||:|:|
Db 67 IHDIIECV 73

RESULT 36
US-11-172-740-1140
;; Sequence 1140, Application US/11172740
;; Publication No. US20060057724A1
;; GENERAL INFORMATION:
;; APPLICANT: MASCIJA, Peter
;; APPLICANT: ALEXANDROV, Nikolai
;; APPLICANT: BROVER, Vyacheslav
;; TITLE OF INVENTION: NUCLEOTIDE SEQUENCES AND POLYPEPTIDES ENCODED THEREBY USEFUL FOR
;; FILE REFERENCE: 2750-1602PUS2
;; CURRENT APPLICATION NUMBER: US/11/172,740
;; CURRENT FILING DATE: 2005-06-30
;; PRIOR APPLICATION NUMBER: 60/583,621
;; PRIOR FILING DATE: 2004-06-30
;; PRIOR APPLICATION NUMBER: 60/584,829
;; PRIOR FILING DATE: 2004-06-30
;; PRIOR APPLICATION NUMBER: 60/584,800
;; PRIOR FILING DATE: 2004-06-30

```

; NUMBER OF SEQ ID NOS: 2523
; SEQ ID NO 1140
; LENGTH: 281
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)-(281)
; OTHER INFORMATION: Ceres CLONE ID no. 463846
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for increasing chlorophyll and photosynthetic cap
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making lethal plants for genetic confinement
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified flower
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making ornamental plants with modified leaves
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with altered leaf shape eg curl
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making plants with increased biomass
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION:
; OTHER INFORMATION: Utility: Useful for making smaller plants
; US-11-172-740-1140

Query Match
Best Local Similarity 37.5%; Score 33; DB 11; Length 281;
Matches 3; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILBCV 8
Db 117 VHDVFIDC 124

RESULT 37
US-11-087-099-9859
; Sequence 9859, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 9859
; LENGTH: 392
; TYPE: PRT
; ORGANISM: Neurospora crassa
; US-11-087-099-9859

Query Match
Best Local Similarity 68.8%; Score 33; DB 11; Length 392;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILBCV 9
Db 87 VYDVHBCV 95
```

```

RESULT 38
US-11-188-298-16877
; Sequence 16877, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16877
; LENGTH: 623
; TYPE: PRT
; ORGANISM: Pichia pastoris
; US-11-188-298-16877

Query Match
Best Local Similarity 62.5%; Score 33; DB 11; Length 623;
Matches 5; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 2 HDIILBCV 9
Db 382 HDIILBCV 389

RESULT 39
US-11-079-463-8639
; Sequence 8639, Application US/11079463
; Publication No. US20060073161A1
; GENERAL INFORMATION:
; APPLICANT: Gary L. Breton
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRA
; FILE REFERENCE: PITH00-03DIY2
; CURRENT APPLICATION NUMBER: US/11/079,463
; CURRENT FILING DATE: 2005-03-14
; PRIOR APPLICATION NUMBER: US 60/128,705
; PRIOR FILING DATE: 1999-04-09
; PRIOR APPLICATION NUMBER: US 09/540,209
; PRIOR FILING DATE: 2000-04-04
; NUMBER OF SEQ ID NOS: 10444
; SEQ ID NO 8639
; LENGTH: 953
; TYPE: PRT
; ORGANISM: B.fragilis
; US-11-079-463-8639

Query Match
Best Local Similarity 71.4%; Score 33; DB 11; Length 953;
Matches 5; Conservative 2; Mismatches 0; Indels 0; Gaps 0;

QY 2 HDIILBCV 8
Db 808 HEMILBCV 814

RESULT 40
US-10-530-253-24
; Sequence 24, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaccia, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
```

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; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 24
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 58
US-10-530-253-24

Query Match      66.7%; Score 32; DB 9; Length 149;
Best Local Similarity 55.6%; Pred. No. 38;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy      1 HDIILECV 9
Db      23 VHEIELKCV 31

RESULT 41
US-10-530-253-25
; Sequence 25, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M17-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 25
; LENGTH: 160
; TYPE: PRT
; ORGANISM: Human papillomavirus type 59
US-10-530-253-25

Query Match      66.7%; Score 32; DB 9; Length 160;
Best Local Similarity 55.6%; Pred. No. 41;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy      1 HDIILECV 9
Db      25 LHDIRINCV 33

RESULT 42
US-11-096-568A-24104
; Sequence 24104, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24104
; LENGTH: 267
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
```

```

; LOCATION: (1)..(267)
; OTHER INFORMATION: Ceres Seq. ID no. 12418821
US-11-096-568A-24104

Query Match      66.7%; Score 32; DB 11; Length 267;
Best Local Similarity 50.0%; Pred. No. 70;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy      2 HDIILECV 9
Db      220 HDCVBECL 227

RESULT 43
US-11-096-568A-24103
; Sequence 24103, Application US/11096568A
; Publication No. US20060048240A1
; GENERAL INFORMATION:
; APPLICANT: Alexandrov, Nikolai et al.
; TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
; FILE REFERENCE: 2750-1592PUS2
; CURRENT APPLICATION NUMBER: US/11/096,568A
; CURRENT FILING DATE: 2005-04-01
; NUMBER OF SEQ ID NOS: 34471
; SEQ ID NO 24103
; LENGTH: 286
; TYPE: PRT
; ORGANISM: Zea mays subsp. mays
; FEATURE:
; NAME/KEY: misc_feature
; LOCATION: (1)..(286)
; OTHER INFORMATION: Ceres Seq. ID no. 12418820
US-11-096-568A-24103

Query Match      66.7%; Score 32; DB 11; Length 286;
Best Local Similarity 50.0%; Pred. No. 75;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy      2 HDIILECV 9
Db      239 HDCVBECL 246

RESULT 44
US-11-087-099-1244
; Sequence 1244, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 1244
; LENGTH: 328
; TYPE: PRT
; ORGANISM: Triticum aestivum
US-11-087-099-1244

Query Match      66.7%; Score 32; DB 11; Length 328;
Best Local Similarity 50.0%; Pred. No. 86;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

Qy      2 HDIILECV 9
Db      220 YDVVEECV 227

RESULT 45
US-11-096-568A-24102
; Sequence 24102, Application US/11096568A
```



```
Publication No. US20060048240A1
GENERAL INFORMATION:
APPLICANT: Alexandrov, Nikolai et al.
TITLE OF INVENTION: Sequence-Determined DNA Fragments and Corresponding Polypeptides
TITLE OF INVENTION: Theby
FILE REFERENCE: 2750-1592PUS2
CURRENT APPLICATION NUMBER: US/11/096,568A
CURRENT FILING DATE: 2005-04-01
NUMBER OF SEQ ID NOS: 34471
SEQ ID NO 24102
LENGTH: 334
TYPE: PRT
ORGANISM: Zea mays subsp. mays
FEATURE:
NAME/KEY: misc feature
LOCATION: (1)-(334)
OTHER INFORMATION: Ceres Seq. ID no. 12418819
US-11-096-568A-24102

Query Match
Best Local Similarity 66.7%; Score 32; DB 11; Length 334;
Best Local 50.0%; Pred. No. 88;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 2 HDIILECV 9
DB 287 HDCEVECL 294

RESULT 46
US-11-188-298-16409
Sequence 16409, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 16409
LENGTH: 340
TYPE: PRT
ORGANISM: Prochlorococcus marinus subsp. pastoris str. CCMPI378
US-11-188-298-16409

Query Match
Best Local Similarity 66.7%; Score 32; DB 11; Length 340;
Best Local 100.0%; Pred. No. 89;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 3 DIIILEC 8
DB 90 DIIILEC 95

RESULT 47
US-11-188-298-646
Sequence 646, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 646
LENGTH: 418
TYPE: PRT
ORGANISM: Burkholderia fungorum
```

```
US-11-188-298-646

Query Match
Best Local Similarity 66.7%; Score 32; DB 11; Length 418;
Best Local 71.4%; Pred. No. 1.1e+02;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 HDIILEC 8
DB 248 HDIILEC 254

RESULT 48
US-11-079-463-5827
Sequence 5827, Application US/11079463
Publication No. US20060073161A1
GENERAL INFORMATION:
APPLICANT: Gary L. Bretton
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO BACTERIOIDES FRAGILIS
TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
FILE REFERENCE: PAT00-03DIV2
CURRENT APPLICATION NUMBER: US/11/079,463
CURRENT FILING DATE: 2005-03-14
PRIOR APPLICATION NUMBER: US 60/128,705
PRIOR FILING DATE: 1999-04-09
PRIOR APPLICATION NUMBER: US 09/540,209
PRIOR FILING DATE: 2000-04-04
NUMBER OF SEQ ID NOS: 10444
SEQ ID NO 5827
LENGTH: 592
TYPE: PRT
ORGANISM: B. fragilis
US-11-079-463-5827

Query Match
Best Local Similarity 66.7%; Score 32; DB 11; Length 592;
Best Local 62.5%; Pred. No. 1.6e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 2 HDIILECV 9
DB 548 HNCIILECV 555

RESULT 49
US-10-505-928-102
Sequence 102, Application US/10505928
Publication No. US20060088532A1
GENERAL INFORMATION:
APPLICANT: Ludwig Institute for Cancer Research et al.
TITLE OF INVENTION: LYMPHATIC ENDOTHELIAL GENES
FILE REFERENCE: 28967/39178
CURRENT APPLICATION NUMBER: US/10/505,928
CURRENT FILING DATE: 2004-08-27
PRIOR APPLICATION NUMBER: US 60/363,019
PRIOR FILING DATE: 2002-03-07
NUMBER OF SEQ ID NOS: 866
SOFTWARE: PatentIn 3.2
SEQ ID NO 102
LENGTH: 780
TYPE: PRT
ORGANISM: Homo sapiens
US-10-505-928-102

Query Match
Best Local Similarity 66.7%; Score 32; DB 8; Length 780;
Best Local 50.0%; Pred. No. 2.1e+02;
Matches 4; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 HDIILEC 8
DB 655 HSLVLDC 662

RESULT 50
US-11-091-668-2
```

```
; Sequence 2, Application US/11091668
; Publication No. US20050262585A1
; GENERAL INFORMATION:
; APPLICANT: University of Nebraska
; APPLICANT: Mackenzie, Sally Ann
; APPLICANT: Vagchindawala, Zarir Erach
; TITLE OF INVENTION: Soybean FGM Synchase Promoters Useful in Parasite Control
; FILE REFERENCE: 1231-221
; CURRENT APPLICATION NUMBER: US/11/091,668
; CURRENT FILING DATE: 2005-03-28
; PRIOR APPLICATION NUMBER: 60556745
; PRIOR FILING DATE: 2004-03-26
; NUMBER OF SEQ ID NOS: 18
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 2
; LENGTH: 1044
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; NAME/KEY: misc feature
; LOCATION: (269)..(269)
; OTHER INFORMATION: The 'Xaa' at location 269 stands for Leu, or Phe.
US-11-091-668-2

Query Match 66.7%; Score 32; DB 11; Length 1044;
Best Local Similarity 55.6%; Pred. No. 2.8e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 IHDIILECV 9
Db 135 VHDRTECV 143

Search completed: May 5, 2006, 08:29:41
Job time : 9.4 secs
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GenCore version 5.1.7
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OM protein - protein search, using sw model

Run on: May 5, 2006, 04:01:20 ; Search time 20.7 Seconds
(Without alignments)
35.946 Million cell updates/sec

Title: US-08-170-344-9
Perfect score: 52
Sequence: 1 CYYCKQQL 9

Scoring table: BLOSUM62
Gapop 10.0, Gapext 0.5

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 572060

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
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Pred. No. is the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	52	100.0	10	2	US-08-159-339A-575
2	52	100.0	20	1	US-08-934-915-160
3	52	100.0	151	2	US-09-701-080C-18
4	52	100.0	158	2	US-09-980-523A-2
5	52	100.0	162	1	US-08-316-239B-3
6	52	100.0	162	1	US-08-316-239B-4
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11	52	100.0	266	2	US-08-860-165-10
12	52	100.0	266	2	US-09-359-382-10
13	52	100.0	266	2	US-09-367-309A-1
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17	52	100.0	390	2	US-09-485-885-14
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22	37	71.2	344	2	US-09-134-000C-5535
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32	35	67.3	383	2	US-09-485-885-23	Sequence 23, Appl
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36	34	65.4	9	2	US-08-159-339A-238	Sequence 238, App
37	34	65.4	9	2	US-08-159-339A-253	Sequence 253, App
38	34	65.4	9	2	US-08-159-339A-570	Sequence 570, App
39	34	65.4	10	2	US-08-159-339A-573	Sequence 573, App
40	34	65.4	15	2	US-08-159-339A-1176	Sequence 1176, Ap
41	34	65.4	32	1	US-08-466-285-2	Sequence 2, Appl1
42	34	65.4	32	2	US-08-164-768-2	Sequence 2, Appl1
43	34	65.4	64	2	US-09-732-210-488	Sequence 488, App
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55	34	65.4	536	2	US-09-977-261-12	Sequence 12, Appl
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57	34	65.4	557	2	US-09-138-277C-3	Sequence 3, Appl1
58	34	65.4	1007	2	US-10-144-198-36	Sequence 36, Appl
59	34	65.4	1041	2	US-10-144-198-14	Sequence 14, Appl
60	34	65.4	3730	2	US-09-949-016-9908	Sequence 9908, Ap
61	33	63.5	17	2	US-09-128-344A-21	Sequence 21, Appl
62	33	63.5	17	2	US-09-128-344A-109	Sequence 109, App
63	33	63.5	17	2	US-09-128-344A-130	Sequence 130, App
64	33	63.5	17	5	US-10-255-011-21	Sequence 21, Appl
65	33	63.5	17	5	US-10-255-011-130	Sequence 130, App
66	33	63.5	17	5	US-10-255-011-109	Sequence 109, App
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69	33	63.5	53	2	US-09-382-911-33	Sequence 33, Appl
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82	33	63.5	375	2	US-10-011-749-22	Sequence 22, Appl
83	33	63.5	383	2	US-09-134-000C-4139	Sequence 4139, Ap
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87	33	63.5	465	2	US-10-011-749-24	Sequence 24, Appl
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91	33	63.5	538	2	US-09-309-572-12	Sequence 12, Appl
92	33	63.5	538	2	US-09-718-096-11	Sequence 11, Appl
93	33	63.5	560	2	US-09-949-016-8096	Sequence 8096, Ap
94	33	63.5	567	1	US-08-841-483-2	Sequence 2, Appl1
95	33	63.5	567	2	US-09-382-911-2	Sequence 2, Appl1
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97	33	63.5	1587	2	US-09-000-094-46	Sequence 46, Appl
98	33	63.5	1587	2	US-10-011-749-46	Sequence 46, Appl
99	33	63.5	1737	2	US-09-309-572-13	Sequence 13, Appl
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102	32	61.5	18	2	US-09-128-344A-71	Sequence 71, Appl	175	31	59.6	196	2	US-09-621-011-174	Sequence 174, App
103	32	61.5	18	2	US-10-255-011-71	Sequence 71, Appl	176	31	59.6	202	2	US-09-248-796A-2499	Sequence 2499, App
104	32	61.5	60	2	US-09-270-767-58063	Sequence 58063, A	177	31	59.6	207	2	US-09-489-039A-11516	Sequence 11516, A
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106	32	61.5	64	2	US-09-621-976-5985	Sequence 5985, Ap	179	31	59.6	292	2	US-09-205-586-667	Sequence 667, App
107	32	61.5	72	2	US-09-248-796A-23292	Sequence 23292, A	180	31	59.6	301	2	US-10-004-860-667	Sequence 667, App
108	32	61.5	76	2	US-09-248-796A-27024	Sequence 27024, A	181	31	59.6	301	2	US-09-148-545-232	Sequence 232, App
109	32	61.5	80	2	US-10-101-464A-985	Sequence 985, App	182	31	59.6	301	2	US-09-621-011-232	Sequence 232, App
110	32	61.5	101	2	US-09-489-039A-7696	Sequence 7696, Ap	183	31	59.6	317	2	US-10-012-331A-322	Sequence 322, App
111	32	61.5	109	2	US-09-248-796A-19570	Sequence 19570, A	184	31	59.6	317	2	US-10-015-389A-322	Sequence 322, App
112	32	61.5	110	2	US-08-545-809A-129	Sequence 129, App	185	31	59.6	317	2	US-10-006-768A-322	Sequence 322, App
113	32	61.5	110	2	US-09-515-697-129	Sequence 129, App	186	31	59.6	317	2	US-10-015-671A-322	Sequence 322, App
114	32	61.5	134	2	US-09-248-796A-19660	Sequence 19660, A	187	31	59.6	317	2	US-10-011-833A-322	Sequence 322, App
115	32	61.5	138	2	US-09-134-001C-4650	Sequence 4650, Ap	188	31	59.6	317	2	US-10-006-041A-322	Sequence 322, App
116	32	61.5	149	2	US-09-270-767-40624	Sequence 40624, A	189	31	59.6	317	2	US-10-012-064A-322	Sequence 322, App
117	32	61.5	153	2	US-09-270-767-55840	Sequence 55840, A	190	31	59.6	327	2	US-09-489-039A-9084	Sequence 9084, Ap
118	32	61.5	153	2	US-09-270-767-41674	Sequence 41674, A	191	31	59.6	334	2	US-09-205-258-672	Sequence 672, App
119	32	61.5	198	2	US-09-328-352-5264	Sequence 5264, Ap	192	31	59.6	334	2	US-10-004-860-672	Sequence 672, App
120	32	61.5	221	2	US-09-248-796A-23414	Sequence 23414, A	193	31	59.6	335	2	US-10-012-331A-130	Sequence 130, App
121	32	61.5	263	2	US-09-662-254B-15	Sequence 15, Appl	194	31	59.6	335	2	US-10-015-389A-130	Sequence 130, App
122	32	61.5	274	2	US-09-538-092-669	Sequence 669, App	195	31	59.6	335	2	US-10-006-768A-130	Sequence 130, App
123	32	61.5	290	2	US-09-538-092-669	Sequence 669, App	196	31	59.6	335	2	US-10-006-768A-130	Sequence 130, App
124	32	61.5	500	2	US-09-328-352-6895	Sequence 6895, Ap	197	31	59.6	335	2	US-10-015-393A-130	Sequence 130, App
125	32	61.5	578	2	US-09-999-248A-2	Sequence 2, Appl	198	31	59.6	335	2	US-10-011-833A-130	Sequence 130, App
126	32	61.5	578	2	US-10-418-036-8	Sequence 8, Appl	199	31	59.6	335	2	US-10-006-041A-130	Sequence 130, App
127	32	61.5	629	2	US-09-198-452A-843	Sequence 843, App	200	31	59.6	335	2	US-10-012-064A-130	Sequence 130, App
128	32	61.5	640	2	US-09-438-185A-795	Sequence 795, App	201	31	59.6	335	2	US-09-134-001C-3672	Sequence 3672, App
129	32	61.5	667	2	US-09-710-279-198	Sequence 198, App	202	31	59.6	399	2	US-09-252-991A-26424	Sequence 26424, A
130	32	61.5	784	2	US-09-740-235-1	Sequence 1, Appl	203	31	59.6	408	2	US-09-902-540-9891	Sequence 6, Appl
131	32	61.5	789	2	US-09-949-016-7164	Sequence 7164, Ap	204	31	59.6	415	2	US-09-006-353A-6	Sequence 6, Appl
132	32	61.5	835	2	US-09-248-796A-15074	Sequence 15074, A	205	31	59.6	415	2	US-09-573-986-6	Sequence 6, Appl
133	32	61.5	886	2	US-09-487-558B-180	Sequence 180, App	206	31	59.6	415	2	US-09-949-016-10234	Sequence 10234, A
134	32	61.5	895	2	US-09-270-767-42746	Sequence 42746, A	207	31	59.6	432	2	US-09-832-129-41	Sequence 41, Appl
135	32	61.5	989	2	US-09-248-796A-19109	Sequence 19109, A	208	31	59.6	432	2	US-09-832-129-54	Sequence 54, Appl
136	32	61.5	1257	2	US-09-920-641-3	Sequence 3, Appl	209	31	59.6	437	2	US-09-252-991A-24572	Sequence 24572, A
137	32	61.5	1500	2	US-09-949-016-5878	Sequence 5878, App	210	31	59.6	466	2	US-09-832-129-59	Sequence 59, Appl
138	31.5	60.6	594	2	US-09-252-991A-20345	Sequence 20345, A	211	31	59.6	466	2	US-08-904-452-2	Sequence 2, Appl
139	31	59.6	18	2	US-09-128-344A-29	Sequence 29, Appl	212	31	59.6	486	2	US-09-517-639-2	Sequence 2, Appl
140	31	59.6	18	2	US-09-128-344A-39	Sequence 39, Appl	213	31	59.6	486	2	US-08-426-009A-15	Sequence 15, Appl
141	31	59.6	18	2	US-09-128-344A-43	Sequence 43, Appl	214	31	59.6	529	2	US-08-232-545-15	Sequence 15, Appl
142	31	59.6	18	2	US-09-128-344A-113	Sequence 113, App	215	31	59.6	529	2	US-09-538-092-885	Sequence 885, App
143	31	59.6	18	2	US-09-128-344A-118	Sequence 118, App	216	31	59.6	529	2	US-09-977-261-15	Sequence 15, Appl
144	31	59.6	18	2	US-09-128-344A-120	Sequence 120, App	217	31	59.6	529	4	PCT-US95-05008-15	Sequence 15, Appl
145	31	59.6	18	2	US-09-128-344A-134	Sequence 134, App	218	31	59.6	529	4	US-09-270-767-61572	Sequence 61572, A
146	31	59.6	18	2	US-09-128-344A-139	Sequence 139, App	219	31	59.6	598	2	US-09-248-796A-18678	Sequence 18678, A
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148	31	59.6	18	5	US-10-255-011-39	Sequence 39, Appl	221	31	59.6	668	2	US-09-270-767-61572	Sequence 18678, A
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153	31	59.6	18	5	US-10-255-011-120	Sequence 120, App	226	31	59.6	721	2	US-10-104-047-2378	Sequence 2378, App
154	31	59.6	18	5	US-10-255-011-134	Sequence 134, App	227	31	59.6	795	2	US-09-489-039A-8249	Sequence 8249, App
155	31	59.6	18	5	US-10-255-011-139	Sequence 139, App	228	31	59.6	808	2	US-08-904-452-4	Sequence 4, Appl
156	31	59.6	18	5	US-10-255-011-141	Sequence 141, App	229	31	59.6	894	2	US-09-517-639-4	Sequence 4, Appl
157	31	59.6	18	5	US-09-270-767-62274	Sequence 62274, A	230	31	59.6	894	2	US-10-060-332-4	Sequence 4, Appl
158	31	59.6	56	2	US-09-205-258-1193	Sequence 1193, App	231	31	59.6	894	2	US-10-339-657-4	Sequence 4, Appl
159	31	59.6	56	2	US-10-004-860-1193	Sequence 1193, App	232	31	59.6	894	2	US-10-885-879-4	Sequence 4, Appl
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161	31	59.6	61	2	US-09-513-999C-498	Sequence 6498, Ap	234	31	59.6	973	2	US-08-904-452-4	Sequence 4, Appl
162	31	59.6	63	2	US-09-277-431A-3	Sequence 3, Appl	235	31	59.6	1068	2	US-09-248-796A-16119	Sequence 16119, A
163	31	59.6	78	2	US-09-448-806C-6	Sequence 6, Appl	236	31	59.6	1085	2	US-09-095-881-2	Sequence 2, Appl
164	31	59.6	83	2	US-09-270-767-48343	Sequence 48343, A	237	31	59.6	1095	2	US-09-555-554-4	Sequence 4, Appl
165	31	59.6	102	2	US-09-540-236-2162	Sequence 2162, Ap	238	31	59.6	1380	2	US-09-949-016-11688	Sequence 11688, A
166	31	59.6	117	2	US-09-543-681A-6635	Sequence 6635, Ap	239	31	59.6	14	1	US-08-771-602D-33	Sequence 33, Appl
167	31	59.6	137	2	US-09-913-204-17	Sequence 17, Appl	240	30	57.7	14	1	US-09-254-776B-64	Sequence 64, Appl
168	31	59.6	141	2	US-09-248-796A-21153	Sequence 21153, A	241	30	57.7	16	2	US-09-128-344A-25	Sequence 25, Appl
169	31	59.6	151	2	US-09-286-529-4	Sequence 4, Appl	242	30	57.7	16	2	US-09-128-344A-131	Sequence 131, App
170	31	59.6	166	2	US-09-270-767-49127	Sequence 49127, A	243	30	57.7	16	2	US-09-128-344A-112	Sequence 112, App
171	31	59.6	170	2	US-09-205-258-1189	Sequence 1189, Ap	244	30	57.7	16	5	US-10-255-011-25	Sequence 25, Appl
172	31	59.6	170	2	US-10-004-860-1189	Sequence 1189, Ap	245	30	57.7	16	5	US-10-255-011-111	Sequence 111, App
173	31	59.6	194	2	US-09-710-279-140	Sequence 140, App	246	30	57.7	16	5	US-10-255-011-111	Sequence 111, App

247	30	57.7	16	5	US-10-255-011-132	Sequence 132, App	320	30	57.7	18	5	US-10-255-011-125	Sequence 125, App
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250	30	57.7	17	2	US-09-128-344A-47	Sequence 47, App1	323	30	57.7	18	5	US-10-255-011-137	Sequence 137, App
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252	30	57.7	17	2	US-09-128-344A-95	Sequence 95, App1	325	30	57.7	18	5	US-10-255-011-144	Sequence 144, App
253	30	57.7	17	2	US-09-128-344A-105	Sequence 105, App	326	30	57.7	18	5	US-10-255-011-198	Sequence 198, App
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257	30	57.7	17	2	US-09-128-344A-131	Sequence 131, App	330	30	57.7	47	2	US-09-270-767-45030	Sequence 60519, A
258	30	57.7	17	2	US-09-128-344A-131	Sequence 143, App	331	30	57.7	47	2	US-09-270-767-60519	Sequence 6, App11
259	30	57.7	17	2	US-08-928-074-19	Sequence 19, App1	332	30	57.7	61	2	US-09-277-431A-4	Sequence 6, App11
260	30	57.7	17	5	US-10-255-011-13	Sequence 13, App1	333	30	57.7	64	2	US-09-762-960-6	Sequence 6, App11
261	30	57.7	17	5	US-10-255-011-23	Sequence 23, App1	334	30	57.7	64	2	US-10-662-756A-6	Sequence 14, App1
262	30	57.7	17	5	US-10-255-011-47	Sequence 47, App1	335	30	57.7	67	1	US-08-511-485-14	Sequence 14, App1
263	30	57.7	17	5	US-10-255-011-93	Sequence 93, App1	336	30	57.7	67	2	US-09-201-936-14	Sequence 14, App1
264	30	57.7	17	5	US-10-255-011-95	Sequence 95, App1	337	30	57.7	67	2	US-09-011-356-14	Sequence 14, App1
265	30	57.7	17	5	US-10-255-011-105	Sequence 105, App	338	30	57.7	67	2	US-09-201-932-14	Sequence 14, App1
266	30	57.7	17	5	US-10-255-011-110	Sequence 110, App	339	30	57.7	67	2	US-09-513-999C-4483	Sequence 4483, App
267	30	57.7	17	5	US-10-255-011-122	Sequence 122, App	340	30	57.7	76	2	US-09-270-767-58747	Sequence 58747, A
268	30	57.7	17	5	US-10-255-011-126	Sequence 126, App	341	30	57.7	76	2	US-09-134-001C-3402	Sequence 3402, App
269	30	57.7	17	5	US-10-255-011-131	Sequence 131, App	342	30	57.7	91	2	US-09-134-001C-3402	Sequence 1064, App
270	30	57.7	17	5	US-10-255-011-143	Sequence 143, App	343	30	57.7	94	2	US-09-732-210-1064	Sequence 5962, A
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283	30	57.7	18	2	US-09-128-344A-86	Sequence 86, App1	356	30	57.7	155	2	US-08-583-110-4800	Sequence 2800, App
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294	30	57.7	18	2	US-09-128-344A-127	Sequence 127, App	367	30	57.7	179	1	US-08-688-342-3	Sequence 3, App11
295	30	57.7	18	2	US-09-128-344A-135	Sequence 135, App	368	30	57.7	179	1	US-09-113-788-3	Sequence 9, App11
296	30	57.7	18	2	US-09-128-344A-137	Sequence 137, App	369	30	57.7	179	2	US-09-113-789-9	Sequence 130, App
297	30	57.7	18	2	US-09-128-344A-142	Sequence 142, App	370	30	57.7	179	2	US-09-919-039-130	Sequence 6200, App
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307	30	57.7	18	5	US-10-255-011-35	Sequence 35, App1	380	30	57.7	227	6	5451506-3	Patent No. 5451506
308	30	57.7	18	5	US-10-255-011-45	Sequence 45, App1	381	30	57.7	227	6	5451506-2	Patent No. 5451506
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313	30	57.7	18	5	US-10-255-011-100	Sequence 100, App	386	30	57.7	252	2	US-08-988-819-13	Sequence 13, App1
314	30	57.7	18	5	US-10-255-011-104	Sequence 104, App	387	30	57.7	252	2	US-09-016-534-13	Sequence 13, App1
315	30	57.7	18	5	US-10-255-011-106	Sequence 106, App	388	30	57.7	252	2	US-08-097-869-3	Sequence 3, App11
316	30	57.7	18	5	US-10-255-011-114	Sequence 114, App	389	30	57.7	252	2	US-09-462-941-21	Sequence 21, App1
317	30	57.7	18	5	US-10-255-011-116	Sequence 116, App	390	30	57.7	254	2	US-09-276-147B-12	Sequence 12, App1
318	30	57.7	18	5	US-10-255-011-121	Sequence 121, App	391	30	57.7	254	2	US-09-270-767-53405	Sequence 53405, A
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394	30	57.7	273	2	US-09-248-796A-15062	Sequence 15062, A	467	29	55.8	25	2	US-09-288-143-94	Sequence 94, Appl
395	30	57.7	275	1	US-08-511-485-11	Sequence 11, Appl	468	29	55.8	26	1	US-08-620-151-125	Sequence 125, App
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398	30	57.7	275	2	US-09-201-936-12	Sequence 12, Appl	471	29	55.8	47	2	US-10-144-929-121	Sequence 121, App
399	30	57.7	275	2	US-09-489-7372	Sequence 7372, Ap	472	29	55.8	47	2	US-09-716-536-14	Sequence 14, Appl
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401	30	57.7	275	2	US-09-201-932-12	Sequence 12, Appl	474	29	55.8	52	2	US-09-513-996C-6575	Sequence 59743, A
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432	30	57.7	470	2	US-09-333-075-6	Sequence 6, Appl1	505	29	55.8	117	2	US-09-270-767-46782	Sequence 880, App
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465	30	57.7	1102	2	US-09-364-609-8	Sequence 8, Appl1	538	29	55.8	178	2	US-10-288-273-6	Sequence 62406, A

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540	29	55.8	188	2	US-09-198-452A-183	Sequence 183, App	613	29	55.8	401	2	US-08-577-788C-4	Sequence 4, Appli
541	29	55.8	188	2	US-09-248-796A-18895	Sequence 18895, A	614	29	55.8	401	2	US-08-577-788C-54	Sequence 54, Appli
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543	29	55.8	199	2	US-09-270-767-45196	Sequence 45196, A	616	29	55.8	411	2	US-09-605-702B-2112	Sequence 2012, Ap
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547	29	55.8	208	2	US-08-577-788C-50	Sequence 50, Appli	620	29	55.8	422	1	US-07-817-920-3	Sequence 3, Appli
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549	29	55.8	212	2	US-09-270-767-44947	Sequence 44947, A	622	29	55.8	422	1	US-08-117-006-3	Sequence 3, Appli
550	29	55.8	219	1	US-08-796-676-1	Sequence 1, Appli	623	29	55.8	422	1	US-08-216-594-3	Sequence 3, Appli
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552	29	55.8	222	2	US-09-710-279-1510	Sequence 1510, Ap	625	29	55.8	422	2	US-08-157-185-13	Sequence 13, Appli
553	29	55.8	222	2	US-09-248-796A-18894	Sequence 18894, A	626	29	55.8	422	2	US-08-281-526B-13	Sequence 3, Appli
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557	29	55.8	247	2	US-09-543-681A-5522	Sequence 5522, Ap	630	29	55.8	438	2	US-09-268-544B-36	Sequence 2, Appli
558	29	55.8	247	2	US-09-134-001C-3569	Sequence 3569, Ap	631	29	55.8	438	2	US-09-950-902-2	Sequence 2, Appli
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832	28	53.8	18	5	US-10-255-011-4	Sequence 4, Appli	905	28	53.8	131	2	US-09-621-976-7697	Sequence 7697, Ap
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853	28	53.8	45	2	US-10-138-618-12	Sequence 12, Appl	926	28	53.8	169	2	US-09-540-236-2860	Sequence 2860, Ap
854	28	53.8	45	2	US-09-690-825-12	Sequence 12, Appl	927	28	53.8	169	2	US-09-398-412B-15	Sequence 15, Appl
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860	28	53.8	54	2	US-09-658-400-6	Sequence 6, Appli	933	28	53.8	180	2	US-09-270-767-92142	Sequence 32142, A
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872	28	53.8	66	2	US-09-511-485-11	Sequence 11, Appl	945	28	53.8	193	2	US-08-818-655-9	Sequence 9, Appli
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874	28	53.8	67	2	US-09-011-356-11	Sequence 11, Appl	947	28	53.8	193	2	US-09-305-839-9	Sequence 9, Appli
875	28	53.8	67	2	US-09-201-932-11	Sequence 11, Appl	948	28	53.8	193	2	US-09-305-839-10	Sequence 10, Appl
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989 28 53.8 331 2 US-09-543-681A-8103 Sequence 8103, A
990 28 53.8 332 2 US-09-134-001C-3977 Sequence 3977, A
991 28 53.8 335 2 US-09-489-039A-10919 Sequence 10919, A
992 28 53.8 338 2 US-09-328-352-6824 Sequence 6824, A
993 28 53.8 345 2 US-09-182-145-3 Sequence 3, Appl 1
994 28 53.8 345 2 US-09-182-145-5 Sequence 5, Appl 1
995 28 53.8 345 2 US-09-182-145-6 Sequence 6, Appl 1
996 28 53.8 345 2 US-09-182-145-11 Sequence 11, Appl 1
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998 28 53.8 347 2 US-09-270-767-45788 Sequence 45788, A
999 28 53.8 349 2 US-09-328-352-5364 Sequence 5364, A
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ALIGNMENTS

RESULT 1
US-08-159-339A-575
; Sequence 575, Application US/08159339A
; Patent No. 6037135
; GENERAL INFORMATION:
; APPLICANT: Kubo, Ralph T.
; APPLICANT: Grey, Howard M.
; APPLICANT: Sette, Alessandro
; APPLICANT: Celis, Esteban
; TITLE OF INVENTION: HLA Binding peptides and Their
; NUMBER OF SEQUENCES: 1254
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Townsend and Townsend and Crew LLP
; STREET: Two Embarcadero Center, Eighth Floor
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111-3834
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Diskette
; COMPUTER: IBM Compatible
; OPERATING SYSTEM: DOS
; SOFTWARE: Pasteo for Windows Version 2.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/159,339A
; FILING DATE: 29-NOV-1993
; CLASSIFICATION: 424
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: US 07/926,666
; FILING DATE: 07-AUG-1992
; APPLICATION NUMBER: US 08/027,746
; FILING DATE: 05-MAR-1993
; APPLICATION NUMBER: US 08/103,396
; FILING DATE: 06-AUG-1993
; ATTORNEY/AGENT INFORMATION:
; NAME: Weber, Ellen Lauver
; REGISTRATION NUMBER: 32,762
; REFERENCE/DOCKET NUMBER: 018623-005030US
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (415) 576-0200
; TELEFAX: (415) 576-0300
; TELEX:
; INFORMATION FOR SEQ ID NO: 575:

SEQUENCE CHARACTERISTICS:
; LENGTH: 10 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-159-339A-575

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Best Local Similarity 100.0%; Pred. No. 0.0073;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 1 CVYCKQQL 9
Db 1 CVYCKQQL 9

RESULT 2
US-08-934-915-160
; Sequence 160, Application US/08934915
; Patent No. 5932412
; GENERAL INFORMATION:
; APPLICANT: DILLNER, JOAKIM
; APPLICANT: DILLNER, LENA
; APPLICANT: CHENG, HWEI-MING
; TITLE OF INVENTION: SYNTHETIC PEPTIDES OF HUMAN
; TITLE OF INVENTION: PAPILLOMAVIRUS 1', 5', 6', 8',
; TITLE OF INVENTION: 11, 16, 18, 31, 33 AND 56,
; TITLE OF INVENTION: USEFUL IN IMMUNOASSAY FOR
; NUMBER OF SEQUENCES: 193
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: MASON & ASSOCIATES, P.A.
; STREET: 17757 U.S. HWY. 19 NORTH, SUITE 500
; CITY: CLEARWATER
; STATE: FLORIDA
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: Windows 3.0
; SOFTWARE: Microsoft Word 6.0
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/934,915
; FILING DATE: 22-SEP-1997
; CLASSIFICATION: 435
; PRIOR APPLICATION DATA:
; APPLICATION NUMBER: 07/949,836
; FILING DATE:
; ATTORNEY/AGENT INFORMATION:
; NAME: LOUISE A. FOUTCH
; REGISTRATION NUMBER: 37,133
; REFERENCE/DOCKET NUMBER: 1946.6
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 813-538-3800
; TELEFAX: 813-538-3820
; TELEX:
; INFORMATION FOR SEQ ID NO: 160:

SEQUENCE CHARACTERISTICS:
; LENGTH: 20 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-934-915-160

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Best Local Similarity 100.0%; Pred. No. 0.014;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Cy 1 CVYCKQQL 9
Db 6 CVYCKQQL 14

RESULT 3
US-09-701-080C-18
; Sequence 18, Application US/09701080C
; Patent No. 6864054
; GENERAL INFORMATION:
; APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
; TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 R
; FILE REFERENCE: N73477C GCM
; CURRENT APPLICATION NUMBER: US/09/701,080C
; CURRENT FILING DATE: 2001-02-27
; PRIOR APPLICATION NUMBER: GB 9811303.8
; PRIOR FILING DATE: 1998-05-26
; PRIOR APPLICATION NUMBER: GB 9900157.0
; PRIOR FILING DATE: 1999-01-05
; NUMBER OF SEQ ID NOS: 36
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 18
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-09-701-080C-18

Query Match 100.0%; Score 52; DB 2; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.09;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 30 CYYCKQOLL 38

RESULT 4
US-09-980-523A-2
; Sequence 2, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTOPIC PROTEIN FRAGMENTS OF THE B6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: MOB1 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; CURRENT FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-2

Query Match 100.0%; Score 52; DB 2; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.094;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

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DB 37 CYYCKQOLL 45

RESULT 5
US-08-316-239B-3
; Sequence 3, Application US/08316239B

Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESS: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 817-9453
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 3:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULAR TYPE: Protein
; HYPOTHETICAL: NO
US-08-316-239B-3

Query Match 100.0%; Score 52; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.096;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CYYCKQOLL 9
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DB 37 CYYCKQOLL 45

RESULT 6
US-08-316-239B-4
; Sequence 4, Application US/08316239B
; Patent No. 5679509
; GENERAL INFORMATION:
; APPLICANT: Wheeler, Cosette M.
; APPLICANT: Parmenter, Cheryl A.
; TITLE OF INVENTION: Methods and a diagnostic Aid for
; TITLE OF INVENTION: Distinguishing a Subset of HPV that is Associated with an
; TITLE OF INVENTION: Increased Risk of Developing Cervical Dysplasia and
; TITLE OF INVENTION: Cervical Cancer
; NUMBER OF SEQUENCES: 4
; CORRESPONDENCE ADDRESS:
; ADDRESS: Jagtiani & Associates
; STREET: 6126 Rocky Way Court
; CITY: Centreville
; STATE: VA
; COUNTRY: USA
; ZIP: 20120-3400
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible

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; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/316,239B
; FILING DATE: 30-SEP-1994
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Jagtiani, Ajay A.
; REGISTRATION NUMBER: 35,205
; REFERENCE/DOCKET NUMBER: UNME-0001
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (703) 803-9387
; TELEFAX: (703) 803-9387
; INFORMATION FOR SEQ ID NO: 4:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 162 amino acids
; TYPE: amino acid
; STRANDEDNESS: not relevant
; TOPOLOGY: not relevant
; MOLECULE TYPE: protein
; HYPOTHEICAL: NO
;
US-08-316-239B-4
;
Query Match 100.0%; Score 52; DB 1; Length 162;
Best Local Similarity 100.0%; Pred. No. 0.096;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
Db 37 CVYCKQQL 45

RESULT 7
US-08-860-165-14
; Sequence 14, Application US/08860165A
; Patent No. 6004557
; GENERAL INFORMATION:
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 11227/130
; CURRENT APPLICATION NUMBER: US/08/860,165A
; CURRENT FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 15
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURES:
; OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
;
US-08-860-165-14
;
Query Match 100.0%; Score 52; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
Db 106 CVYCKQQL 114

RESULT 8
US-09-359-382-14
; Sequence 14, Application US/09359382
; Patent No. 6306397
; GENERAL INFORMATION:
```

```
; APPLICANT: EDWARDS, Stirling John
; APPLICANT: COX, John Cooper
; APPLICANT: WEBB, Elizabeth Ann
; APPLICANT: FRAZER, Ian
; TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
; FILE REFERENCE: 017227/0148
; CURRENT APPLICATION NUMBER: US/09/359,382
; CURRENT FILING DATE: 1999-07-23
; EARLIER APPLICATION NUMBER: US 08/860,165
; EARLIER FILING DATE: 1997-09-22
; EARLIER APPLICATION NUMBER: PCT/AU95/00868
; EARLIER FILING DATE: 1995-12-20
; EARLIER APPLICATION NUMBER: AU PNO157/94
; EARLIER FILING DATE: 1994-12-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 14
; LENGTH: 172
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
;
US-09-359-382-14
;
Query Match 100.0%; Score 52; DB 2; Length 172;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
Db 106 CVYCKQQL 114

RESULT 9
US-08-117-083-10
; Sequence 10, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Boursnell, Michael E.
; APPLICANT: Inglish, Stephen C.
; APPLICANT: Munro, Alan J.
; TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
; NUMBER OF SEQUENCES: 70
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Walter H. Dregger
; STREET: 4 Embarcadero Center, Suite 3400
; CITY: San Francisco
; STATE: CA
; COUNTRY: USA
; ZIP: 94111
; COMPUTER READABLE FORM:
; MEDIUM TYPE: floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: Patentin Release #1.0, Version #1.25
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/117,083
; FILING DATE: 10-SEP-1993
; CLASSIFICATION: 435
; ATTORNEY/AGENT INFORMATION:
; NAME: Dregger, Walter H.
; REGISTRATION NUMBER: 24,190
; REFERENCE/DOCKET NUMBER: A-56783
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 415-781-1989
; TELEFAX: 415-398-3249
; TELEX: 910 277299
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 182 amino acids
; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
; MOLECULE TYPE: protein
```

FEATURE:
NAME/KEY: Protein
LOCATION: 1..182
OTHER INFORMATION: /note= "Xaa refers to stop codon in
OTHER INFORMATION: the open reading frame."
US-08-117-083-10

Query Match 100.0%; Score 52; DB 1; Length 182;
Best Local Similarity 100.0%; Pred. No. 0.11;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
DB 38 CVYCKQQL 46

RESULT 10
US-09-462-993-1

Sequence 1, Application US/09462993
Patent No. 6884786
GENERAL INFORMATION:
APPLICANT: KIRBY, Marie-Paule
APPLICANT: BIZOUARNE, Nadine
TITLE OF INVENTION: ANTITUMORAL COMPOSITION BASED ON IMMUNOGENIC
FILE REFERENCE: 01753-122
CURRENT APPLICATION NUMBER: US/09/462,993
CURRENT FILING DATE: 2000-04-17
PRIOR APPLICATION NUMBER: PCT/FR98/01576
PRIOR FILING DATE: 1998-07-17
PRIOR APPLICATION NUMBER: FR 97/09152
PRIOR FILING DATE: 1997-07-18
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.2
SEQ ID NO 1
LENGTH: 243
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURES:
OTHER INFORMATION: Description of Artificial Sequence: Derived from
OTHER INFORMATION: human papillomavirus, strain HPV-16, B6 protein
OTHER INFORMATION: fused F protein signals, clone B6*TMF.
US-09-462-993-1

Query Match 100.0%; Score 52; DB 2; Length 243;
Best Local Similarity 100.0%; Pred. No. 0.14;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
DB 65 CVYCKQQL 73

RESULT 11
US-08-860-165-10

Sequence 10, Application US/08860165A
Patent No. 6004557
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 17227/130
CURRENT APPLICATION NUMBER: US/08/860,165A
CURRENT FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P010157
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 15
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURES:
OTHER INFORMATION: Description of Artificial Sequence: Gene Fusion
US-08-860-165-10

Query Match 100.0%; Score 52; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
DB 37 CVYCKQQL 45

RESULT 12

US-09-359-382-10
Sequence 10, Application US/09359382
Patent No. 6306397
GENERAL INFORMATION:
APPLICANT: EDWARDS, Stirling John
APPLICANT: COX, John Cooper
APPLICANT: WEBB, Elizabeth Ann
APPLICANT: FAZER, Ian
TITLE OF INVENTION: VARIANTS OF HUMAN PAPILLOMA VIRUS ANTIGENS
FILE REFERENCE: 017227/0148
CURRENT APPLICATION NUMBER: US/09/359,382
CURRENT FILING DATE: 1999-07-23
EARLIER APPLICATION NUMBER: US 08/860,165
EARLIER FILING DATE: 1997-09-22
EARLIER APPLICATION NUMBER: PCT/AU95/00868
EARLIER FILING DATE: 1995-12-20
EARLIER APPLICATION NUMBER: AU P010157/94
EARLIER FILING DATE: 1994-12-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 266
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-09-359-382-10

Query Match 100.0%; Score 52; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
DB 37 CVYCKQQL 45

RESULT 13

US-09-367-309A-1
Sequence 1, Application US/09367309A
Patent No. 6428807
GENERAL INFORMATION:
APPLICANT: MACFARLAN, RODERICK I.
APPLICANT: MALLIAROS, JIM
TITLE OF INVENTION: CHELATING IMMUNOSTIMULATING COMPLEXES
FILE REFERENCE: 017227/0149
CURRENT APPLICATION NUMBER: US/09/367,309A
CURRENT FILING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: PCT/AU98/00080
PRIOR FILING DATE: 1998-02-13
PRIOR APPLICATION NUMBER: AU PO 5178
PRIOR FILING DATE: 1997-02-19
NUMBER OF SEQ ID NOS: 6
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 1
LENGTH: 266
TYPE: PRT

ORGANISM: Human papillomavirus type 16
US-09-367-309A-1

Query Match 100.0%; Score 52; DB 2; Length 266;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
DB 37 CVYCKOQL 45

RESULT 14
US-09-485-885-4
Sequence 4, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 4
LENGTH: 273
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-4

Query Match 100.0%; Score 52; DB 2; Length 273;
Best Local Similarity 100.0%; Pred. No. 0.15;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
DB 143 CVYCKOQL 151

RESULT 15
US-09-485-885-10
Sequence 10, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 10
LENGTH: 292
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-10

Query Match 100.0%; Score 52; DB 2; Length 292;
Best Local Similarity 100.0%; Pred. No. 0.16;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
DB 162 CVYCKOQL 170

RESULT 16
US-09-485-885-6
Sequence 6, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 6
LENGTH: 371
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-6

Query Match 100.0%; Score 52; DB 2; Length 371;
Best Local Similarity 100.0%; Pred. No. 0.21;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
DB 143 CVYCKOQL 151

RESULT 17
US-09-485-885-14
Sequence 14, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FastSeq for Windows Version 3.0
SEQ ID NO 14
LENGTH: 390
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-14

Query Match 100.0%; Score 52; DB 2; Length 390;
Best Local Similarity 100.0%; Pred. No. 0.22;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 162 CVYCKQQL 170

RESULT 18
US-09-980-523A-4
; Sequence 4, Application US/09980523A
; Patent No. 6783763
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIER, ESTELLE
; TITLE OF INVENTION: POLYPEPTIC PROTEIN FRAGMENTS OF THE E6 AND E7
; TITLE OF INVENTION: PROTEINS OF HPV, THEIR PRODUCTION AND THEIR USE
; TITLE OF INVENTION: PARTICULARLY IN VACCINATION
; FILE REFERENCE: MOB1 AO INS
; CURRENT APPLICATION NUMBER: US/09/980,523A
; PRIOR FILING DATE: 2002-04-29
; PRIOR APPLICATION NUMBER: PCT/FR00/01513
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: FR 99/07012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: Patent In Ver. 2.1
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Human Papillomavirus
US-09-980-523A-4

Query Match 92.3%; Score 48; DB 2; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 8
|||
Db 23 CVYCKQQL 30

RESULT 19
US-09-252-991A-31270
; Sequence 31270, Application US/09252991A
; Patent No. 6551795
; GENERAL INFORMATION:
; APPLICANT: Marc J. Rubenfield et al.
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO PSEUDOMONAS
; TITLE OF INVENTION: AERUGINOSA FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.136
; CURRENT APPLICATION NUMBER: US/09/252,991A
; CURRENT FILING DATE: 1999-02-18
; PRIOR APPLICATION NUMBER: US 60/074,788
; PRIOR FILING DATE: 1998-02-18
; PRIOR APPLICATION NUMBER: US 60/094,190
; PRIOR FILING DATE: 1998-07-27
; NUMBER OF SEQ ID NOS: 33142
; SEQ ID NO 31270
; LENGTH: 80
; TYPE: PRT
; ORGANISM: Pseudomonas aeruginosa
US-09-252-991A-31270

Query Match 78.8%; Score 41; DB 2; Length 80;
Best Local Similarity 66.7%; Pred. No. 3.9;
Matches 6; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 63 CVYCKQQL 71

RESULT 20
US-09-248-796A-24285
; Sequence 24285, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICA
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 24285
; LENGTH: 347
; TYPE: PRT
; ORGANISM: Candida albicans
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (75)
; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unk.
US-09-248-796A-24285

Query Match 78.8%; Score 41; DB 2; Length 347;
Best Local Similarity 66.7%; Pred. No. 15;
Matches 6; Conservative 1; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 19 CVYCKQQL 27

RESULT 21
US-09-134-001C-4260
; Sequence 4260, Application US/09134001C
; Patent No. 6380370
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO STAPHYLOCOCCU
; TITLE OF INVENTION: EPIDERMIDIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: GTC-007
; CURRENT APPLICATION NUMBER: US/09/134,001C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/064,964
; PRIOR FILING DATE: 1997-11-08
; PRIOR APPLICATION NUMBER: US 60/055,779
; PRIOR FILING DATE: 1997-08-14
; NUMBER OF SEQ ID NOS: 5674
; SEQ ID NO 4260
; LENGTH: 156
; TYPE: PRT
; ORGANISM: Staphylococcus epidermidis
US-09-134-001C-4260

Query Match 73.1%; Score 38; DB 2; Length 156;
Best Local Similarity 44.4%; Pred. No. 23;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 122 CVYCKQQL 130

RESULT 22
US-09-134-000C-5935
; Sequence 5935, Application US/09134000C
; Patent No. 6617156
; GENERAL INFORMATION:
; APPLICANT: Lynn Doucette-Stamm et al

```

; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO
; TITLE OF INVENTION: ENTEROCOCCUS FAECALIS FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 032796-032
; CURRENT APPLICATION NUMBER: US/09/134,000C
; CURRENT FILING DATE: 1998-08-13
; PRIOR APPLICATION NUMBER: US 60/055,778
; PRIOR FILING DATE: 1997-08-15
; NUMBER OF SEQ ID NOS: 6812
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5935
; LENGTH: 344
; TYPE: PRT
; ORGANISM: Enterococcus faecalis
; US-09-134-000C-5935

Query Match          71.2%; Score 37; DB 2; Length 344;
Best Local Similarity 55.6%; Pred. No. 72;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY      1 CVCCKOQL 9
       |||||:
Db      7 CWCCKTQII 15

RESULT 23
US-09-270-767-39036
; Sequence 39036, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 39036
; LENGTH: 959
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
; US-09-270-767-39036

Query Match          71.2%; Score 37; DB 2; Length 959;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 CVCCKOQL 8
       |||||:
Db      732 CXYCKKQI 739

RESULT 24
US-09-270-767-54253
; Sequence 54253, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 54253
; LENGTH: 959
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
; US-09-270-767-54253
```

```

Query Match          71.2%; Score 37; DB 2; Length 959;
Best Local Similarity 62.5%; Pred. No. 1.9e+02;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 CVCCKOQL 8
       |||||:
Db      732 CXYCKKQI 739

RESULT 25
US-10-101-464A-984
; Sequence 984, Application US/10101464A
; Patent No. 6768041
; GENERAL INFORMATION:
; APPLICANT: Strabala, Timothy
; APPLICANT: Nieuwenhuizen, Nicolaas
; APPLICANT: Higgins, Colleen M.
; TITLE OF INVENTION: Compositions Isolated from Plant Cells
; TITLE OF INVENTION: and Their Use in the Modification of Plant Cell Signaling
; FILE REFERENCE: 11000.1020c2
; CURRENT APPLICATION NUMBER: US/10/101,464A
; CURRENT FILING DATE: 2002-03-18
; PRIOR APPLICATION NUMBER: 09/704,302
; PRIOR FILING DATE: 2000-11-01
; PRIOR APPLICATION NUMBER: 09/228,986
; PRIOR FILING DATE: 1999-01-12
; PRIOR APPLICATION NUMBER: 60/162,866
; PRIOR FILING DATE: 1999-11-01
; PRIOR APPLICATION NUMBER: PCT/US00/00724
; PRIOR FILING DATE: 2000-01-11
; NUMBER OF SEQ ID NOS: 989
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 984
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Pinus radiata
; US-10-101-464A-984

Query Match          69.2%; Score 36; DB 2; Length 151;
Best Local Similarity 62.5%; Pred. No. 50;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;

QY      1 CVCCKOQL 8
       |||||:
Db      112 CVCYCLOQM 119

RESULT 26
US-09-949-016-9659
; Sequence 9659, Application US/09949016
; Patent No. 6812339
; GENERAL INFORMATION:
; APPLICANT: VENTER, J. Craig et al.
; TITLE OF INVENTION: POLYMORPHISMS IN KNOWN GENES ASSOCIATED
; TITLE OF INVENTION: WITH HUMAN DISEASE, METHODS OF DETECTION AND USES THEREOF
; FILE REFERENCE: CU001307
; CURRENT APPLICATION NUMBER: US/09/949,016
; CURRENT FILING DATE: 2000-04-14
; PRIOR APPLICATION NUMBER: 60/241,755
; PRIOR FILING DATE: 2000-10-20
; PRIOR APPLICATION NUMBER: 60/237,768
; PRIOR FILING DATE: 2000-10-03
; PRIOR APPLICATION NUMBER: 60/231,498
; PRIOR FILING DATE: 2000-09-08
; NUMBER OF SEQ ID NOS: 207012
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 9659
; LENGTH: 441
; TYPE: PRT
; ORGANISM: Human
; US-09-949-016-9659

Query Match          69.2%; Score 36; DB 2; Length 441;
```


Best Local Similarity 75.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

Qy 1 CVYCKQOL 8
Db 162 CVYCKQOL 169

RESULT 27

US-09-489-039A-12427
; Sequence 12427, Application US/09489039A
; Patent No. 6610836
; GENERAL INFORMATION:
; APPLICANT: Gary Breton et. al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO KLEBSIELLA
; TITLE OF INVENTION: PNEUMONIAE FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 2709.2004001
; CURRENT APPLICATION NUMBER: US/09/489,039A
; CURRENT FILING DATE: 2000-01-27
; PRIOR APPLICATION NUMBER: US 60/117,747
; PRIOR FILING DATE: 1999-01-29
; NUMBER OF SEQ ID NOS: 14342
; SEQ ID NO 12427
; LENGTH: 462
; TYPE: PRT
; ORGANISM: Klebsiella pneumoniae
US-09-489-039A-12427

Query Match 69.2%; Score 36; DB 2; Length 462;
Best Local Similarity 55.6%; Pred. No. 1.4e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKQOL 9
Db 308 CVYCKQOL 316

RESULT 28

US-08-247-904B-10
; Sequence 10, Application US/08247904B
; Patent No. 5981699
; GENERAL INFORMATION:
; APPLICANT: Rolfe, Mark
; APPLICANT: Eckstein, Jens W.
; APPLICANT: Draetta, Giulio
; TITLE OF INVENTION: Human Ubiquitin Conjugating Enzyme
; NUMBER OF SEQUENCES: 17
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: Foley, Hoag & Elliot
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: ASCII(text)
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/247,904B
; FILING DATE: 23-MAY-1994
; CLASSIFICATION: 530
; ATTORNEY/AGENT INFORMATION:
; NAME: Vincent, Matthew P.
; REGISTRATION NUMBER: 36,709
; REFERENCE/DOCKET NUMBER: MIV-029, 01
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: (617) 832-1000
; TELEFAX: (617) 832-7000
; INFORMATION FOR SEQ ID NO: 10:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 158 amino acids

; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-247-904B-10

Query Match 67.3%; Score 35; DB 1; Length 158;
Best Local Similarity 75.0%; Pred. No. 78;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKQOL 8
Db 32 CVYCKQOL 39

RESULT 29

US-08-767-942A-19
; Sequence 19, Application US/08767942A
; Patent No. 6068982
; GENERAL INFORMATION:
; APPLICANT: Rolfe, Mark
; APPLICANT: Chin, M. Isabel
; APPLICANT: Berlin, Vivian
; APPLICANT: Damagomez, Veronique
; APPLICANT: Draetta, Giulio
; APPLICANT: Guillaume, Cottarel
; TITLE OF INVENTION: UBIQUITIN CONJUGATING ENZYMES
; NUMBER OF SEQUENCES: 45
; CORRESPONDENCE ADDRESSES:
; ADDRESSEE: FOLEY, HONG & ELLIOT LLP
; STREET: One Post Office Square
; CITY: Boston
; STATE: MA
; COUNTRY: USA
; ZIP: 02109-2170
; COMPUTER READABLE FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; OPERATING SYSTEM: PC-DOS/MS-DOS
; SOFTWARE: PatentIn Release #1.0, Version #1.30
; CURRENT APPLICATION DATA:
; APPLICATION NUMBER: US/08/767,942A
; FILING DATE: 17-DEC-1996
; ATTORNEY/AGENT INFORMATION:
; NAME: Vincent, Matthew P.
; REGISTRATION NUMBER: 36,709
; REFERENCE/DOCKET NUMBER: MIV-029, 04
; TELECOMMUNICATION INFORMATION:
; TELEPHONE: 617-832-1000
; TELEFAX: 617-832-7000
; INFORMATION FOR SEQ ID NO: 19:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 158 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-767-942A-19

Query Match 67.3%; Score 35; DB 2; Length 158;
Best Local Similarity 75.0%; Pred. No. 78;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKQOL 8
Db 32 CVYCKQOL 39

RESULT 30

US-08-117-083-14
; Sequence 14, Application US/08117083
; Patent No. 5719054
; GENERAL INFORMATION:
; APPLICANT: Bournelli, Michael E.
; APPLICANT: Inglis, Stephen C.

APPLICANT: Munro, Alan J.
TITLE OF INVENTION: Recombinant Virus Vectors Encoding Human
TITLE OF INVENTION: Papilloma Virus Proteins
NUMBER OF SEQUENCES: 70
CORRESPONDENCE ADDRESS:
ADDRESSEE: Walter H. Dreger
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,083
FILING DATE: 10-SEP-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-58783
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-781-1989
TELEFAX: 415-398-3249
TELEX: 910 277239
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 271 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: protein
FEATURE:
NAME/KEY: Protein
LOCATION: 1..271
OTHER INFORMATION: /note="Xaa refers to stop codon in
US-08-117-083-14
OTHER INFORMATION: the open reading frame."

Query Match 67.3%; Score 35; DB 1; Length 271;
Best Local Similarity 75.0%; Pred. No. 1.3e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKOOL 8
Db 33 CVYCKTVL 40

RESULT 31
US-09-485-885-21
Sequence 21, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FaestSeq for Windows Version 3.0
SEQ ID NO 21
LENGTH: 278

TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-21

Query Match 67.3%; Score 35; DB 2; Length 278;
Best Local Similarity 75.0%; Pred. No. 1.3e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKOOL 8
Db 143 CVYCKTVL 150

RESULT 32
US-09-485-885-23
Sequence 23, Application US/09485885
Patent No. 6342224
GENERAL INFORMATION:
APPLICANT: Bruck, Claudine
APPLICANT: Cabezon Silva, Teresa
APPLICANT: Delisse, Anne-Marie Eva Fernande
APPLICANT: Gerard, Catherine Marie Ghislaine
APPLICANT: Lombardo-Bencheikh, Angela
TITLE OF INVENTION: Vaccine
FILE REFERENCE: B45107
CURRENT APPLICATION NUMBER: US/09/485,885
CURRENT FILING DATE: 2000-02-18
PRIOR APPLICATION NUMBER: PCT/EP98/05285
PRIOR FILING DATE: 1998-08-17
PRIOR APPLICATION NUMBER: GB 9717953.5
PRIOR FILING DATE: 1997-08-22
NUMBER OF SEQ ID NOS: 23
SOFTWARE: FaestSeq for Windows Version 3.0
SEQ ID NO 23
LENGTH: 383
TYPE: PRT
ORGANISM: Homo sapien
US-09-485-885-23

Query Match 67.3%; Score 35; DB 2; Length 383;
Best Local Similarity 75.0%; Pred. No. 1.8e+02;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKOOL 8
Db 143 CVYCKTVL 150

RESULT 33
US-09-328-352-4537
Sequence 4537, Application US/09328352
Patent No. 6562958
GENERAL INFORMATION:
APPLICANT: Gary L. Breton et al.
TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO ACINETOBACTER
FILE REFERENCE: GTC99-03PA
CURRENT APPLICATION NUMBER: US/09/328,352
CURRENT FILING DATE: 1999-06-04
NUMBER OF SEQ ID NOS: 8252
SEQ ID NO 4537
LENGTH: 472
TYPE: PRT
ORGANISM: Acinetobacter baumannii
US-09-328-352-4537

Query Match 67.3%; Score 35; DB 2; Length 472;
Best Local Similarity 44.4%; Pred. No. 2.1e+02;
Matches 4; Conservative 3; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKOOL 9
Db 314 CIVNSRML 322

RESULT 34
US-09-138-277C-1

Sequence 1, Application US/09138277C

Patent No. 6426403

GENERAL INFORMATION:

APPLICANT: NAKATA, MOTOMI

APPLICANT: NAKANO, HIROYASU

APPLICANT: YAGITA, HIDEO

APPLICANT: OKUMURA, KO

TITLE OF INVENTION: TRAP FAMILY MOLECULES, POLYNUCLEOTIDES ENCODING THEM,

FILE REFERENCE: 007898-0255515

CURRENT APPLICATION NUMBER: US/09/138,277C

PRIOR APPLICATION NUMBER: PCT/JP97/00512

PRIOR FILING DATE: 1997-02-24

PRIOR APPLICATION NUMBER: JP 34674/1996

NUMBER OF SEQ ID NOS: 16

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 1

LENGTH: 558

TYPE: PRT

ORGANISM: Mus sp.

US-09-138-277C-1

Query Match 67.3%; Score 35; DB 2; Length 558;

Best Local Similarity 44.4%; Pred. No. 2.5e+02;

Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Qy 1 CVCCKQQLL 9

Db 168 CVCCKRDIV 176

RESULT 35
US-08-159-339A-226

Sequence 226, Application US/08159339A

Patent No. 6037135

GENERAL INFORMATION:

APPLICANT: Kubo, Ralph T.

APPLICANT: Grey, Howard M.

APPLICANT: Sette, Alessandro

APPLICANT: Celis, Esben

TITLE OF INVENTION: HLA Binding peptides and Their

NUMBER OF SEQUENCES: 1254

CORRESPONDENCE ADDRESS:

ADDRESS: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

STATE: CA

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FastSeq for Windows Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/159,339A

FILING DATE: 29-NOV-1993

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/926,666

FILING DATE: 07-AUG-1992

APPLICATION NUMBER: US 08/027,746

FILING DATE: 05-MAR-1993

APPLICATION NUMBER: US 08/103,396

FILING DATE: 06-AUG-1993

ATTORNEY/AGENT INFORMATION:

NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:

INFORMATION FOR SEQ ID NO: 226:

SEQUENCE CHARACTERISTICS:

LENGTH: 9 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear

MOLECULE TYPE: peptide

US-08-159-339A-226

Query Match 65.4%; Score 34; DB 2; Length 9;

Best Local Similarity 87.5%; Pred. No. 4.6e+05;

Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

Qy 2 VYCKQQLL 9

Db 2 VYCKQQLL 9

RESULT 36
US-08-159-339A-238

Sequence 238, Application US/08159339A

Patent No. 6037135

GENERAL INFORMATION:

APPLICANT: Kubo, Ralph T.

APPLICANT: Grey, Howard M.

APPLICANT: Sette, Alessandro

APPLICANT: Celis, Esben

TITLE OF INVENTION: HLA Binding peptides and Their

NUMBER OF SEQUENCES: 1254

CORRESPONDENCE ADDRESS:

ADDRESS: Townsend and Townsend and Crew LLP

STREET: Two Embarcadero Center, Eighth Floor

CITY: San Francisco

STATE: CA

COUNTRY: USA

ZIP: 94111-3834

COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: IBM Compatible

OPERATING SYSTEM: DOS

SOFTWARE: FastSeq for Windows Version 2.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/159,339A

FILING DATE: 29-NOV-1993

CLASSIFICATION: 424

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/926,666

FILING DATE: 07-AUG-1992

APPLICATION NUMBER: US 08/027,746

FILING DATE: 05-MAR-1993

APPLICATION NUMBER: US 08/103,396

FILING DATE: 06-AUG-1993

ATTORNEY/AGENT INFORMATION:

NAME: Weber, Ellen Lauver

REGISTRATION NUMBER: 32,762

REFERENCE/DOCKET NUMBER: 018623-005030US

TELECOMMUNICATION INFORMATION:

TELEPHONE: (415) 576-0200

TELEFAX: (415) 576-0300

TELEX:

INFORMATION FOR SEQ ID NO: 238:

SEQUENCE CHARACTERISTICS:

LENGTH: 9 amino acids

TYPE: amino acid

STRANDEDNESS: single

TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-238

Query Match 65.4%; Score 34; DB 2; Length 9;
Best Local Similarity 87.5%; Pred. No. 4.6e+05;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 VYCKQQL 9
DB 1 VYAKQQL 8

RESULT 37
US-08-159-339A-253
Sequence 253, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esben
TITLE OF INVENTION: HLA Binding peptides and Their
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 253:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-253

Query Match 65.4%; Score 34; DB 2; Length 9;
Best Local Similarity 87.5%; Pred. No. 4.6e+05;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

QY 2 VYCKQQL 9
DB 1 VYAKQQL 8

RESULT 38
US-08-159-339A-570
Sequence 570, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esben
TITLE OF INVENTION: HLA Binding peptides and Their
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 570:
SEQUENCE CHARACTERISTICS:
LENGTH: 9 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-570

Query Match 65.4%; Score 34; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCK 5
DB 5 CVYCK 9

RESULT 39
US-08-159-339A-573
Sequence 573, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Esben
TITLE OF INVENTION: HLA Binding peptides and Their

Query Match 65.4%; Score 34; DB 2; Length 9;
Best Local Similarity 100.0%; Pred. No. 4.6e+05;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 573:
SEQUENCE CHARACTERISTICS:
LENGTH: 10 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-573

Query Match 65.4%; Score 34; DB 2; Length 10;
Best Local Similarity 100.0%; Pred. No. 9;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCK 5
DB 6 CVYCK 10

RESULT 40
US-08-159-339A-1176
Sequence 1176, Application US/08159339A
Patent No. 6037135
GENERAL INFORMATION:
APPLICANT: Kubo, Ralph T.
APPLICANT: Grey, Howard M.
APPLICANT: Sette, Alessandro
APPLICANT: Celis, Beseban
TITLE OF INVENTION: HLA Binding peptides and Their
NUMBER OF SEQUENCES: 1254
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Eighth Floor
CITY: San Francisco
STATE: CA
COUNTRY: USA
ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS

SOFTWARE: FastSeq for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/159,339A
FILING DATE: 29-NOV-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/926,666
FILING DATE: 07-AUG-1992
APPLICATION NUMBER: US 08/027,746
FILING DATE: 05-MAR-1993
APPLICATION NUMBER: US 08/103,396
FILING DATE: 06-AUG-1993
ATTORNEY/AGENT INFORMATION:
NAME: Weber, Ellen Lauver
REGISTRATION NUMBER: 32,762
REFERENCE/DOCKET NUMBER: 018623-005030US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
TELEX:
INFORMATION FOR SEQ ID NO: 1176:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-159-339A-1176

Query Match 65.4%; Score 34; DB 2; Length 15;
Best Local Similarity 100.0%; Pred. No. 13;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCK 5
DB 11 CVYCK 15

RESULT 41
US-08-466-285-2
Sequence 2, Application US/08466285
Patent No. 5753233
GENERAL INFORMATION:
APPLICANT: Bleul, Conrad
APPLICANT: Giesmann, Latz
APPLICANT: Muller, Martin
TITLE OF INVENTION: Seroreactive Epitopes On Proteins Of
TITLE OF INVENTION: Human Papillomavirus (HPV)18
NUMBER OF SEQUENCES: 7
CORRESPONDENCE ADDRESS:
ADDRESSEE: Finnegan, Henderson, Farabow, Garrett &
ADDRESS: Dunner
STREET: 1300 I Street, N.W., Suite 700
CITY: Washington
STATE: D.C.
COUNTRY: USA
ZIP: 20005-3315
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/466,285
FILING DATE: 06-JUN-1995
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/947,992
FILING DATE: 21-SEP-1992

CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/696,953
FILING DATE: 08-MAY-1991
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: P 40 15 044.5
FILING DATE: 10-MAY-1990
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Manepelzer, David A.
REGISTRATION NUMBER: 37,540
REFERENCE/DOCKET NUMBER: 05552.1075-03000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202)408-4400
TELEFAX: (202)408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-466-285-2

Query Match 65.4%; Score 34; DB 1; Length 32;
Best Local Similarity 100.0%; Pred. No. 26;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCK 5
DB 27 CVCCK 31

RESULT 42
US-08-164-768-2
Sequence 2, Application US/08164768
Patent No. 6322794
GENERAL INFORMATION:
APPLICANT: BLEUL, Conrad
APPLICANT: GISSMANN, Lutz
APPLICANT: MITLER, Martin
TITLE OF INVENTION: SEROREACTIVE EPITOPES ON PROTEINS OF
NUMBER OF SEQUENCES: 7
TITLE OF INVENTION: HUMAN PAPILLOMA VIRUS (HPV) 18
CORRESPONDENCE ADDRESS:
ADDRESSEE: PINNEGAN, HENDERSON, FARABOW, GARRETT &
ADDRESS: DUNNER, L.L.P.
STREET: 1300 I Street, N.W.
CITY: Washington
STATE: DC
COUNTRY: USA
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/164,768
FILING DATE: 10-DEC-1993
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Forman, David S.
REGISTRATION NUMBER: 33,694
REFERENCE/DOCKET NUMBER: 05552.1075-02000
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 408-4000
TELEFAX: (202) 408-4400
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 32 amino acids
TYPE: amino acid

STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
US-08-164-768-2

Query Match 65.4%; Score 34; DB 2; Length 32;
Best Local Similarity 100.0%; Pred. No. 26;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCK 5
DB 27 CVCCK 31

RESULT 43
US-09-732-210-488
Sequence 488, Application US/09732210
Patent No. 6573361
GENERAL INFORMATION:
APPLICANT: Bunkers, Greg J.
APPLICANT: Liang, Jihong
APPLICANT: Miltanck, Cindy A.
APPLICANT: Seale, Jeffrey W.
APPLICANT: Wu, Yonnie S.
TITLE OF INVENTION: Anti-fungal proteins and methods for their use
FILE REFERENCE: 38-21(15036)B
CURRENT APPLICATION NUMBER: US/09/732,210
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: US 60/169,513
PRIOR FILING DATE: 1999-12-07
PRIOR APPLICATION NUMBER: US 60/169,340
PRIOR FILING DATE: 1999-12-07
NUMBER OF SEQ ID NOS: 1753
SEQ ID NO 488
LENGTH: 64
TYPE: PRT
ORGANISM: Cyanophora paradoxa
US-09-732-210-488

Query Match 65.4%; Score 34; DB 2; Length 64;
Best Local Similarity 55.6%; Pred. No. 50;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVCCKOOL 9
DB 51 CVCCKOOL 59

RESULT 44
US-09-270-767-60607
Sequence 60607, Application US/09270767
Patent No. 6703491
GENERAL INFORMATION:
APPLICANT: Homburger et al.
TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
FILE REFERENCE: File Reference: 7326-094
CURRENT APPLICATION NUMBER: US/09/270,767
CURRENT FILING DATE: 1999-03-17
NUMBER OF SEQ ID NOS: 62517
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 60607
LENGTH: 118
TYPE: PRT
ORGANISM: Drosophila melanogaster
FEATURE:
OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-60607

Query Match 65.4%; Score 34; DB 2; Length 118;
Best Local Similarity 83.3%; Pred. No. 88;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCKQ 6

Db 5 CVYCKQ 10

RESULT 45
US-09-248-796A-14459

; Sequence 14459, Application US/09248796A
; Patent No. 6747137
; GENERAL INFORMATION:
; APPLICANT: Keith Weinstock et al
; TITLE OF INVENTION: NUCLEIC ACID AND AMINO ACID SEQUENCES RELATING TO CANDIDA ALBICAN
; TITLE OF INVENTION: FOR DIAGNOSTICS AND THERAPEUTICS
; FILE REFERENCE: 107196.132
; CURRENT APPLICATION NUMBER: US/09/248,796A
; CURRENT FILING DATE: 1999-02-12
; PRIOR APPLICATION NUMBER: US 60/074,725
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: US 60/096,409
; PRIOR FILING DATE: 1998-08-13
; NUMBER OF SEQ ID NOS: 28208
; SEQ ID NO 14459
; LENGTH: 168
; TYPE: PRT
; ORGANISM: Candida albicans
; FEATURE:
; NAME/KEY: UNSURE
; LOCATION: (2)
; OTHER INFORMATION: Identity of amino acid sequences at the above locations are unknc
US-09-248-796A-14459

Query Match 65.4%; Score 34; DB 2; Length 168;
Best Local Similarity 55.6%; Pred. No. 1.2e+02;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
Db 153 CYYMKQQLM 161

RESULT 46
US-09-270-767-40146

; Sequence 40146, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 40146
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-40146

Query Match 65.4%; Score 34; DB 2; Length 273;
Best Local Similarity 71.4%; Pred. No. 1.9e+02;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQ 7
Db 200 CVYCETQ 206

RESULT 47
US-09-270-767-55362
; Sequence 55362, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:

; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094

; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 55362
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:

; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-55362

Query Match 65.4%; Score 34; DB 2; Length 273;
Best Local Similarity 71.4%; Pred. No. 1.9e+02;
Matches 5; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQ 7
Db 200 CVYCETQ 206

RESULT 48
US-09-270-767-37306

; Sequence 37306, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 37306
; LENGTH: 334
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-37306

Query Match 65.4%; Score 34; DB 2; Length 334;
Best Local Similarity 75.0%; Pred. No. 2.3e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 VYCKQQL 9
Db 61 VYCKQQL 68

RESULT 49
US-09-270-767-52523

; Sequence 52523, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of Drosophila melanogaster
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270,767
; CURRENT FILING DATE: 1999-03-17
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: PatentIn Ver. 2.0
; SEQ ID NO 52523
; LENGTH: 334
; TYPE: PRT
; ORGANISM: Drosophila melanogaster
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-52523

Query Match 65.4%; Score 34; DB 2; Length 334;
Best Local Similarity 75.0%; Pred. No. 2.3e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 2 VYCKQQL 9
| | | | |
Db 61 VYCEQQL 68

RESULT 50

US-09-270-767-62069
; Sequence 62069, Application US/09270767
; Patent No. 6703491
; GENERAL INFORMATION:
; APPLICANT: Homburger et al.
; TITLE OF INVENTION: Nucleic acids and proteins of *Drosophila melanogaster*
; FILE REFERENCE: File Reference: 7326-094
; CURRENT APPLICATION NUMBER: US/09/270.767
; NUMBER OF SEQ ID NOS: 62517
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 62069
; LENGTH: 344
; TYPE: PRT
; ORGANISM: *Drosophila melanogaster*
; FEATURE:
; OTHER INFORMATION: Xaa means any amino acid
US-09-270-767-62069

Query Match 65.4%; Score 34; DB 2; Length 344;
Best Local Similarity 44.4%; Pred. No. 2.4e+02;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
| | | | |
Db 238 CIPCKTKVL 246

Search completed: May 5, 2006, 04:48:42
Job time : 22.7 secs

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OM protein - protein search, using sw model

Run on: May 5, 2006, 08:18:14 ; Search time 56 Seconds
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Title: US-08-170-344-9
Perfect score: 52
Sequence: 1 CVYCKQQL 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 1867569 seqs, 417829326 residues
Total number of hits satisfying chosen parameters: 1867569

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database : Published Applications_AA_Main:*

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- 2: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.dep:*
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- 4: /cgn2_6/ptodata/1/pubpaa/US10A_PUBCOMB.dep:*
- 5: /cgn2_6/ptodata/1/pubpaa/US10B_PUBCOMB.dep:*
- 6: /cgn2_6/ptodata/1/pubpaa/US11_PUBCOMB.dep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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2	52	100.0	15	4	US-10-476-570-24
3	52	100.0	15	4	US-10-476-570-25
4	52	100.0	21	4	US-10-476-570-10
5	52	100.0	32	4	US-10-476-570-9
6	52	100.0	33	4	US-10-476-570-19
7	52	100.0	151	4	US-10-177-390-6
8	52	100.0	151	5	US-10-484-063-20
9	52	100.0	151	5	US-10-484-063-27
10	52	100.0	158	5	US-10-858-384-2
11	52	100.0	158	5	US-10-367-057-16
12	52	100.0	158	6	US-11-021-949-13
13	52	100.0	171	4	US-10-472-724-2
14	52	100.0	243	6	US-11-072-288-1
15	52	100.0	266	3	US-09-367-309A-1
16	52	100.0	273	4	US-10-000-903-4
17	52	100.0	273	5	US-10-899-771-4
18	52	100.0	292	4	US-10-899-771-10
19	52	100.0	292	5	US-10-000-903-10
20	52	100.0	371	4	US-10-000-903-6
21	52	100.0	371	5	US-10-899-771-6
22	52	100.0	390	4	US-10-000-903-14
23	52	100.0	390	5	US-10-899-771-14
24	52	100.0	536	4	US-10-367-095-10
25	52	100.0	536	4	US-10-368-046-10
26	52	100.0	536	4	US-10-367-367-10
27	52	100.0	536	5	US-10-918-337-10

28	48	92.3	30	4	US-10-476-570-53	Sequence 53, Appl
29	48	92.3	30	5	US-10-858-384-4	Sequence 4, Appl
30	45	86.5	149	6	US-11-021-949-14	Sequence 14, Appl
31	42	80.8	158	6	US-11-021-949-361	Sequence 361, App
32	41	78.8	149	6	US-11-021-949-18	Sequence 18, Appl
33	41	78.8	151	6	US-11-021-949-24	Sequence 24, Appl
34	41	78.8	151	6	US-11-021-949-25	Sequence 25, Appl
35	41	78.8	153	6	US-11-021-949-20	Sequence 20, Appl
36	41	78.8	155	6	US-11-021-949-22	Sequence 22, Appl
37	41	78.8	155	6	US-11-021-949-23	Sequence 23, Appl
38	41	78.8	162	6	US-11-021-949-31	Sequence 31, Appl
39	40	76.9	135	4	US-10-437-963-106779	Sequence 106779, Sequence 359, App
40	40	76.9	148	6	US-11-021-949-359	Sequence 23, Appl
41	39	75.0	15	4	US-10-476-570-23	Sequence 11069, A
42	39	75.0	147	4	US-10-369-493-11069	Sequence 27, Appl
43	39	75.0	150	6	US-11-021-949-27	Sequence 229548, Sequence 312020,
44	39	75.0	157	4	US-10-424-599-229548	Sequence 6, Appl
45	39	75.0	185	4	US-10-425-115-312020	Sequence 35, Appl
46	39	75.0	398	4	US-10-291-737-6	Sequence 5326, Ap
47	39	75.0	398	4	US-10-365-564-6	Sequence 199640, Sequence 26, Appl
48	39	75.0	449	4	US-10-225-810-35	Sequence 300, App
49	39	75.0	449	4	US-10-369-493-5326	Sequence 18751, A
50	38	73.1	145	4	US-10-425-115-185751	Sequence 159035, Sequence 234065,
51	38	73.1	151	6	US-11-021-949-26	Sequence 145353, Sequence 135, App
52	38	73.1	156	4	US-10-724-972A-3796	Sequence 20331, Sequence 78, Appl
53	38	73.1	343	4	US-10-369-493-300	Sequence 61457, A
54	38	73.1	774	4	US-10-425-114-43519	Sequence 330, App
55	38	73.1	798	4	US-10-425-115-58706	Sequence 11324, A
56	38	73.1	1724	4	US-10-437-963-159035	Sequence 58706, A
57	37	71.2	45	4	US-10-425-115-234065	Sequence 6996, Ap
58	37	71.2	79	4	US-10-437-963-145353	Sequence 6996, Ap
59	37	71.2	88	4	US-10-023-896-135	Sequence 21247, A
60	37	71.2	88	6	US-11-122-117-135	Sequence 18, Appl
61	37	71.2	96	4	US-10-437-963-202331	Sequence 4, Appl
62	37	71.2	107	4	US-10-023-896-78	Sequence 169, App
63	37	71.2	107	6	US-11-122-117-78	Sequence 169006, Sequence 227351,
64	37	71.2	137	4	US-10-767-701-61457	Sequence 34870, A
65	37	71.2	183	3	US-09-895-913A-330	Sequence 223437, A
66	37	71.2	328	3	US-09-815-242-11324	Sequence 369075, A
67	37	71.2	328	4	US-10-282-122A-58706	Sequence 10115, A
68	37	71.2	328	4	US-10-335-977-6595	Sequence 984, App
69	37	71.2	331	4	US-10-335-977-6596	Sequence 984, App
70	37	71.2	344	4	US-10-369-493-21247	Sequence 21247, A
71	37	71.2	1958	4	US-10-467-433-18	Sequence 18, Appl
72	37	71.2	2669	4	US-10-016-248-4	Sequence 4, Appl
73	37	71.2	4823	4	US-10-051-874-169	Sequence 169, App
74	36	69.2	67	4	US-10-424-599-166006	Sequence 227351, A
75	36	69.2	70	4	US-10-425-115-227351	Sequence 34870, A
76	36	69.2	96	4	US-10-767-701-61457	Sequence 223437, A
77	36	69.2	105	4	US-10-425-115-223437	Sequence 369075, A
78	36	69.2	108	4	US-10-425-115-369075	Sequence 10115, A
79	36	69.2	132	5	US-10-739-930-10115	Sequence 984, App
80	36	69.2	151	4	US-10-101-464A-984	Sequence 984, App
81	36	69.2	151	5	US-10-864-252-984	Sequence 29, Appl
82	36	69.2	158	6	US-11-021-949-29	Sequence 34839, A
83	36	69.2	311	4	US-10-425-115-311839	Sequence 41600, A
84	36	69.2	383	5	US-10-450-763-41860	Sequence 10, Appl
85	36	69.2	397	3	US-09-932-367A-10	Sequence 3, Appl
86	36	69.2	399	3	US-09-932-367A-3	Sequence 14, Appl
87	36	69.2	401	3	US-09-932-367A-14	Sequence 12, Appl
88	36	69.2	402	3	US-09-932-367A-12	Sequence 21, Appl
89	36	69.2	402	3	US-09-932-367A-21	Sequence 1959, Ap
90	36	69.2	402	5	US-10-723-860-1959	Sequence 16, Appl
91	36	69.2	403	3	US-09-932-367A-16	Sequence 700, App
92	36	69.2	457	4	US-10-369-493-700	Sequence 13785, A
93	36	69.2	2144	6	US-11-097-143-13785	Sequence 1439, App
94	35	67.3	35	5	US-10-751-845-139	Sequence 1449, Ap
95	35	67.3	35	5	US-10-808-167-1449	Sequence 1449, Ap
96	35	67.3	35	5	US-10-807-807-1449	Sequence 154, App
97	35	67.3	42	5	US-10-751-845-152	Sequence 238732, A
98	35	67.3	87	4	US-10-425-115-238732	Sequence 159, App
99	35	67.3	119	5	US-10-751-845-159	
100	35	67.3				

101	35	67.3	122	4	US-10-425-114-48646	Sequence 48646, A	174	34	65.4	677	5	US-10-450-763-37731	Sequence 37731, A
102	35	67.3	122	4	US-10-425-115-348124	Sequence 348124, A	175	34	65.4	708	5	US-10-450-763-56138	Sequence 56138, A
103	35	67.3	127	3	US-09-925-301-1213	Sequence 1213, Ap	176	34	65.4	708	5	US-10-450-763-58760	Sequence 58760, A
104	35	67.3	158	3	US-10-800-023-27	Sequence 27, Appl	177	34	65.4	721	5	US-10-450-763-37386	Sequence 37386, A
105	35	67.3	158	3	US-11-021-949-28	Sequence 28, Appl	178	34	65.4	725	4	US-10-108-260A-3060	Sequence 3060, Ap
106	35	67.3	158	6	US-11-021-949-30	Sequence 30, Appl	179	34	65.4	747	6	US-11-097-143-39840	Sequence 39840, A
107	35	67.3	172	4	US-10-472-724-6	Sequence 6, Appl1	180	34	65.4	795	5	US-10-450-763-37384	Sequence 37384, A
108	35	67.3	236	5	US-10-751-845-157	Sequence 157, App	181	34	65.4	820	5	US-10-450-763-38762	Sequence 38762, A
109	35	67.3	237	5	US-10-751-845-158	Sequence 158, App	182	34	65.4	824	4	US-10-437-963-171380	Sequence 171380, A
110	35	67.3	261	5	US-10-751-845-160	Sequence 160, App	183	34	65.4	841	5	US-10-450-763-38766	Sequence 38767, A
111	35	67.3	278	5	US-10-000-903-21	Sequence 21, Appl	184	34	65.4	890	4	US-10-437-963-171378	Sequence 37866, A
112	35	67.3	278	5	US-10-899-771-21	Sequence 21, Appl	185	34	65.4	892	4	US-10-437-963-171372	Sequence 171372, A
113	35	67.3	343	4	US-10-282-122A-60799	Sequence 60799, A	186	34	65.4	948	6	US-11-097-143-957	Sequence 957, App
114	35	67.3	383	4	US-10-000-903-23	Sequence 23, Appl	187	34	65.4	965	4	US-10-437-963-171385	Sequence 171385, A
115	35	67.3	383	5	US-10-899-771-23	Sequence 23, Appl	188	34	65.4	984	6	US-11-097-143-37704	Sequence 37704, A
116	35	67.3	394	5	US-09-932-367A-5	Sequence 5, Appl1	189	34	65.4	1002	3	US-09-812-671-3	Sequence 3, Appl1
117	35	67.3	395	3	US-09-932-367A-4	Sequence 4, Appl1	190	34	65.4	1002	3	US-09-812-633-3	Sequence 3, Appl1
118	35	67.3	398	3	US-09-932-367A-6	Sequence 6, Appl1	191	34	65.4	1002	3	US-09-988-117-3	Sequence 3, Appl1
119	35	67.3	422	6	US-11-097-143-18057	Sequence 18057, A	192	34	65.4	1002	3	US-10-765-520-3	Sequence 3, Appl1
120	35	67.3	558	4	US-10-004-378A-32	Sequence 32, Appl	193	34	65.4	1002	5	US-10-828-815-3	Sequence 3, Appl1
121	35	67.3	558	4	US-10-004-378A-33	Sequence 33, Appl	194	34	65.4	1002	5	US-10-828-815-3	Sequence 3, Appl1
122	35	67.3	558	4	US-10-042-865-164	Sequence 164, App	195	34	65.4	1005	3	US-09-812-471-1	Sequence 1, Appl1
123	35	67.3	558	4	US-10-042-865-165	Sequence 165, App	196	34	65.4	1005	3	US-09-812-633-1	Sequence 1, Appl1
124	35	67.3	1096	4	US-10-473-576-3	Sequence 3, Appl1	197	34	65.4	1005	3	US-09-988-117-1	Sequence 1, Appl1
125	35	67.3	1119	4	US-10-170-385-13	Sequence 13, Appl	198	34	65.4	1005	4	US-10-197-824-29	Sequence 29, Appl
126	35	67.3	1139	4	US-11-097-143-10833	Sequence 10833, A	199	34	65.4	1005	4	US-10-765-520-1	Sequence 1, Appl1
127	34	65.4	53	4	US-10-424-599-205112	Sequence 205112, A	200	34	65.4	1005	5	US-10-828-815-1	Sequence 1, Appl1
128	34	65.4	53	4	US-10-087-887-71	Sequence 71, Appl	201	34	65.4	1007	4	US-10-144-198-36	Sequence 36, Appl
129	34	65.4	60	3	US-09-864-761-37174	Sequence 37174, A	202	34	65.4	1007	5	US-10-197-824-38	Sequence 38, Appl
130	34	65.4	61	4	US-10-087-887-82	Sequence 82, Appl	203	34	65.4	1007	5	US-10-486-977-24	Sequence 24, Appl
131	34	65.4	73	4	US-10-437-963-166617	Sequence 166617, A	204	34	65.4	103	4	US-10-437-963-171373	Sequence 171373, A
132	34	65.4	86	3	US-10-425-115-353924	Sequence 353924, A	205	34	65.4	1041	4	US-10-144-198-14	Sequence 14, Appl
133	34	65.4	86	3	US-09-764-868-916	Sequence 916, App	206	34	65.4	1118	4	US-10-437-963-125420	Sequence 125420, A
134	34	65.4	87	4	US-10-767-701-50513	Sequence 50513, A	207	34	65.4	1350	4	US-10-437-963-125423	Sequence 125423, A
135	34	65.4	88	4	US-10-425-115-185638	Sequence 185638, A	208	34	65.4	1780	4	US-10-437-963-11376	Sequence 17176, A
136	34	65.4	88	4	US-10-425-115-164657	Sequence 264657, A	209	34	65.4	3572	4	US-10-471-450-16	Sequence 16, Appl
137	34	65.4	93	3	US-09-764-868-1223	Sequence 1223, Ap	210	34	65.4	17	3	US-09-865-943-109	Sequence 21, Appl
138	34	65.4	96	3	US-09-864-761-43066	Sequence 43066, A	211	33	63.5	17	3	US-09-865-943-130	Sequence 109, App
139	34	65.4	100	4	US-10-424-599-274744	Sequence 274744, A	212	33	63.5	28	6	US-11-077-752-17	Sequence 170, App
140	34	65.4	100	4	US-10-425-115-359834	Sequence 359834, A	213	33	63.5	74	4	US-10-424-599-223973	Sequence 223973, A
141	34	65.4	105	4	US-10-425-115-338126	Sequence 338126, A	214	33	63.5	76	4	US-10-425-115-193939	Sequence 190399, A
142	34	65.4	119	4	US-10-767-701-32284	Sequence 32284, A	215	33	63.5	77	4	US-10-425-115-328497	Sequence 328497, A
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145	34	65.4	165	4	US-10-425-115-341522	Sequence 341522, A	218	33	63.5	101	4	US-10-424-599-185597	Sequence 185597, A
146	34	65.4	190	4	US-10-437-963-171381	Sequence 171381, A	219	33	63.5	101	4	US-10-424-599-247626	Sequence 247626, A
147	34	65.4	190	4	US-10-437-963-178593	Sequence 178593, A	220	33	63.5	102	4	US-10-437-963-155784	Sequence 155784, A
148	34	65.4	247	4	US-10-425-115-192033	Sequence 192033, A	221	33	63.5	114	4	US-10-425-115-239490	Sequence 239490, A
149	34	65.4	297	4	US-10-424-599-195734	Sequence 195734, A	222	33	63.5	134	4	US-10-425-115-355127	Sequence 355127, A
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152	34	65.4	320	4	US-09-939-853A-11	Sequence 11, Appl	225	33	63.5	152	4	US-10-425-115-354973	Sequence 34973, A
153	34	65.4	320	4	US-10-087-887-12	Sequence 12, Appl	226	33	63.5	166	4	US-10-425-115-356865	Sequence 356190, A
154	34	65.4	323	4	US-10-425-114-69151	Sequence 69151, A	227	33	63.5	173	4	US-10-425-115-366865	Sequence 366865, A
155	34	65.4	329	4	US-10-363-829-317	Sequence 317, App	228	33	63.5	183	4	US-10-437-963-131915	Sequence 131915, A
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158	34	65.4	526	4	US-10-262-445-130	Sequence 130, App	231	33	63.5	197	4	US-10-108-260A-3343	Sequence 3343, Ap
159	34	65.4	526	4	US-10-042-865-50	Sequence 50, Appl	232	33	63.5	203	4	US-10-437-963-138264	Sequence 138264, A
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161	34	65.4	536	3	US-09-977-261-12	Sequence 12, Appl	234	33	63.5	228	4	US-10-406-686A-56	Sequence 56, Appl
162	34	65.4	536	5	US-10-732-923-13649	Sequence 13649, A	235	33	63.5	228	4	US-10-425-114-67396	Sequence 67396, A
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164	34	65.4	557	4	US-10-004-378A-34	Sequence 34, Appl	237	33	63.5	242	4	US-10-264-213-248	Sequence 248, App
165	34	65.4	557	4	US-10-042-865-162	Sequence 162, App	238	33	63.5	251	4	US-10-437-963-138213	Sequence 138213, A
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167	34	65.4	559	4	US-10-087-887-14	Sequence 14, Appl	240	33	63.5	384	4	US-10-437-963-133671	Sequence 132671, A
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172	34	65.4	574	6	US-11-097-143-40230	Sequence 40230, A	245	33	63.5				
173	34	65.4	660	5	US-10-741-849-7144	Sequence 7144, Ap	246	33	63.5				

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249	33	63.5	403	3	US-09-805-456-6	Sequence 6, Appl1	322	32	61.5	114	4	US-10-372-766-160	Sequence 160, App
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258	33	63.5	508	4	US-10-282-122A-42524	Sequence 53574, A	331	32	61.5	145	4	US-10-372-876-427	Sequence 876-427
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262	33	63.5	567	4	US-10-369-022-2	Sequence 24, Appl1	335	32	61.5	162	4	US-10-437-963-137223	Sequence 137223,
263	33	63.5	567	5	US-10-480-068-24	Sequence 25, Appl1	336	32	61.5	162	4	US-10-799-747-296	Sequence 296, App
264	33	63.5	567	5	US-10-480-068-25	Sequence 52648, A	337	32	61.5	162	4	US-10-799-747-296	Sequence 296, App
265	33	63.5	580	5	US-10-450-763-52648	Sequence 32, Appl1	338	32	61.5	178	5	US-10-450-763-32471	Sequence 32471, A
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267	33	63.5	625	4	US-10-408-765A-2049	Sequence 863, App	340	32	61.5	190	4	US-10-437-963-176230	Sequence 176230,
268	33	63.5	670	5	US-10-631-467-863	Sequence 863, App	341	32	61.5	200	5	US-10-450-763-38330	Sequence 38330, A
269	33	63.5	690	5	US-10-631-467-862	Sequence 236, App	342	32	61.5	200	5	US-10-450-763-38330	Sequence 38330, A
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271	33	63.5	693	4	US-10-437-963-177190	Sequence 39750, A	344	32	61.5	214	3	US-09-782-980-23	Sequence 980-23
272	33	63.5	696	6	US-11-097-143-39750	Sequence 324, App	345	32	61.5	214	4	US-10-303-502-2	Sequence 502-2
273	33	63.5	734	4	US-10-408-765A-324	Sequence 297, App	346	32	61.5	222	4	US-10-806-018-23	Sequence 806-018-23
274	33	63.5	734	5	US-10-723-860-297	Sequence 2050, Ap	347	32	61.5	224	4	US-10-425-115-117438	Sequence 117438
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278	33	63.5	779	4	US-10-437-963-194924	Sequence 48603, A	351	32	61.5	248	4	US-10-767-701-10668	Sequence 10668, A
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284	33	63.5	1846	6	US-11-097-143-16281	Sequence 53682, A	357	32	61.5	263	5	US-10-450-763-40162	Sequence 40162, A
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287	33	63.5	5809	5	US-10-450-763-53654	Sequence 53654, A	360	32	61.5	298	4	US-10-282-122A-61140	Sequence 61140
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293	32	61.5	31	4	US-10-799-747-302	Sequence 302, App	366	32	61.5	334	5	US-10-276-774-2054	Sequence 774-2054
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297	32	61.5	49	4	US-10-425-115-280319	Sequence 280319, A	370	32	61.5	346	4	US-10-723-860-3318	Sequence 723-860-3318
298	32	61.5	50	4	US-10-437-963-180292	Sequence 180292, A	371	32	61.5	346	5	US-10-654-102-92	Sequence 654-102-92
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305	32	61.5	68	4	US-10-437-963-109291	Sequence 109291, A	378	32	61.5	411	4	US-10-238-075-11102	Sequence 238-075-11102
306	32	61.5	68	4	US-10-425-115-306231	Sequence 306231, A	379	32	61.5	416	3	US-09-780-532-6	Sequence 780-532-6
307	32	61.5	70	4	US-10-424-599-183381	Sequence 183381, A	380	32	61.5	417	3	US-10-052-586-474	Sequence 586-474
308	32	61.5	70	4	US-10-767-701-50060	Sequence 50060, A	381	32	61.5	417	4	US-10-176-550-474	Sequence 176-550-474
309	32	61.5	77	4	US-10-264-049-3159	Sequence 3159, Ap	382	32	61.5	417	4	US-10-176-758-474	Sequence 758-474
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312	32	61.5	82	4	US-10-767-701-58509	Sequence 203311, A	385	32	61.5	417	4	US-10-176-483-474	Sequence 483-474
313	32	61.5	84	4	US-10-424-599-203311	Sequence 203311, A	386	32	61.5	417	4	US-10-176-799-474	Sequence 799-474
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316	32	61.5	98	4	US-10-424-599-227337	Sequence 227337, A	389	32	61.5	417	4	US-10-173-706-474	Sequence 706-474
317	32	61.5	98	4	US-10-437-963-133500	Sequence 133500, A	390	32	61.5	417	4	US-10-175-728-474	Sequence 728-474
318	32	61.5	102	4	US-10-425-115-331215	Sequence 331215, A	391	32	61.5	417	4	US-10-175-752-474	Sequence 752-474
319	32	61.5	108	3	US-09-925-300-982	Sequence 982, App	392	32	61.5	417	4	US-10-175-752-474	Sequence 752-474

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394	32	61.5	417	4	US-10-176-913-474	Sequence 414, App	467	32	61.5	417	4	US-10-180-913-474	Sequence 474, App
395	32	61.5	417	4	US-10-176-517-474	Sequence 414, App	468	32	61.5	417	4	US-10-184-612-474	Sequence 474, App
396	32	61.5	417	4	US-10-180-552-474	Sequence 414, App	469	32	61.5	417	4	US-10-184-616-474	Sequence 474, App
397	32	61.5	417	4	US-10-180-557-474	Sequence 414, App	470	32	61.5	417	4	US-10-184-617-474	Sequence 474, App
398	32	61.5	417	4	US-10-173-700-474	Sequence 414, App	471	32	61.5	417	4	US-10-184-622-474	Sequence 474, App
399	32	61.5	417	4	US-10-174-572-474	Sequence 414, App	472	32	61.5	417	4	US-10-184-628-474	Sequence 474, App
400	32	61.5	417	4	US-10-174-579-474	Sequence 414, App	473	32	61.5	417	4	US-10-184-629-474	Sequence 474, App
401	32	61.5	417	4	US-10-174-582-474	Sequence 414, App	474	32	61.5	417	4	US-10-184-631-474	Sequence 474, App
402	32	61.5	417	4	US-10-174-588-474	Sequence 414, App	475	32	61.5	417	4	US-10-184-632-474	Sequence 474, App
403	32	61.5	417	4	US-10-175-739-474	Sequence 414, App	476	32	61.5	417	4	US-10-184-636-474	Sequence 474, App
404	32	61.5	417	4	US-10-175-740-474	Sequence 414, App	477	32	61.5	417	4	US-10-184-640-474	Sequence 474, App
405	32	61.5	417	4	US-10-175-743-474	Sequence 414, App	478	32	61.5	417	4	US-10-184-650-474	Sequence 474, App
406	32	61.5	417	4	US-10-176-488-474	Sequence 414, App	479	32	61.5	417	4	US-10-184-651-474	Sequence 474, App
407	32	61.5	417	4	US-10-176-492-474	Sequence 414, App	480	32	61.5	417	4	US-10-184-652-474	Sequence 474, App
408	32	61.5	417	4	US-10-176-747-474	Sequence 414, App	481	32	61.5	417	4	US-10-187-568-474	Sequence 474, App
409	32	61.5	417	4	US-10-176-747-474	Sequence 414, App	482	32	61.5	417	4	US-10-187-597-474	Sequence 474, App
410	32	61.5	417	4	US-10-176-885-474	Sequence 414, App	483	32	61.5	417	4	US-10-187-598-474	Sequence 474, App
411	32	61.5	417	4	US-10-176-887-474	Sequence 414, App	484	32	61.5	417	4	US-10-187-600-474	Sequence 474, App
412	32	61.5	417	4	US-10-176-992-474	Sequence 414, App	485	32	61.5	417	4	US-10-187-601-474	Sequence 474, App
413	32	61.5	417	4	US-10-176-993-474	Sequence 414, App	486	32	61.5	417	4	US-10-187-603-474	Sequence 474, App
414	32	61.5	417	4	US-10-184-658-474	Sequence 414, App	487	32	61.5	417	4	US-10-187-606-474	Sequence 474, App
415	32	61.5	417	4	US-10-176-991-474	Sequence 414, App	488	32	61.5	417	4	US-10-187-741-474	Sequence 474, App
416	32	61.5	417	4	US-10-227-884-220	Sequence 220, App	489	32	61.5	417	4	US-10-187-743-474	Sequence 474, App
417	32	61.5	417	4	US-10-173-695-474	Sequence 414, App	490	32	61.5	417	4	US-10-187-746-474	Sequence 474, App
418	32	61.5	417	4	US-10-173-697-474	Sequence 414, App	491	32	61.5	417	4	US-10-187-747-474	Sequence 474, App
419	32	61.5	417	4	US-10-173-705-474	Sequence 414, App	492	32	61.5	417	4	US-10-187-751-474	Sequence 474, App
420	32	61.5	417	4	US-10-174-576-474	Sequence 414, App	493	32	61.5	417	4	US-10-187-753-474	Sequence 474, App
421	32	61.5	417	4	US-10-174-585-474	Sequence 414, App	494	32	61.5	417	4	US-10-187-754-474	Sequence 474, App
422	32	61.5	417	4	US-10-174-586-474	Sequence 414, App	495	32	61.5	417	4	US-10-187-757-474	Sequence 474, App
423	32	61.5	417	4	US-10-175-747-474	Sequence 414, App	496	32	61.5	417	4	US-10-187-884-474	Sequence 474, App
424	32	61.5	417	4	US-10-175-747-474	Sequence 414, App	497	32	61.5	417	4	US-10-188-767-474	Sequence 474, App
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ALIGNMENTS

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US-10-484-063-3
; Sequence 3, Application US/10484063
; Publication No. US20050048467A1
; GENERAL INFORMATION:
; APPLICANT: SASSTRY, K. JACANNADRA
; APPLICANT: TORTOLERO-LINA, GUILLEMO
; APPLICANT: FOLLEN, MICHELE
; TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
; FILE REFERENCE: UISC:560US
; CURRENT APPLICATION NUMBER: US/10/484, 063
; CURRENT FILING DATE: 2004-01-16
; PRIOR APPLICATION NUMBER: PCT/US02/23198
; PRIOR FILING DATE: 2002-07-19
; PRIOR APPLICATION NUMBER: 60/306, 809
; PRIOR FILING DATE: 2001-07-20
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 3
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-484-063-3

Query Match 100.0%; Score 52; DB 5; Length 10;
Best Local Similarity 100.0%; Pred. No. 0.06;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476, 570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 24
; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
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; OTHER INFORMATION: Description of the artificial sequence: peptide E6 31-45
US-10-476-570-24

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Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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; Publication No. US20040170644A1
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; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORATILLE, Sandra
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476, 570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
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RESULT 4
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; Publication No. US20040170644A1
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APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 10
LENGTH: 21
TYPE: PRT
ORGANISM: artificial sequence
FEATURES:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 30-50
US-10-476-570-10
```

```
Query Match          100.0%; Score 52; DB 4; Length 21;
Best Local Similarity 100.0%; Pred. No. 0.12;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 8 CVYCKQQL 16
```

```
RESULT 5
US-10-476-570-9
Sequence 9, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 9
LENGTH: 32
TYPE: PRT
ORGANISM: artificial sequence
FEATURES:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-45
US-10-476-570-9
```

```
Query Match          100.0%; Score 52; DB 4; Length 32;
Best Local Similarity 100.0%; Pred. No. 0.17;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 24 CVYCKQQL 32
```

```
RESULT 6
US-10-476-570-19
Sequence 19, Application US/10476570
Publication No. US20040170644A1
GENERAL INFORMATION:
APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
APPLICANT: MAILLERE, Bernard
APPLICANT: BOURGAULT-VILLADA, Isabelle
APPLICANT: POUVELLE-MORATILLE, Sandra
APPLICANT: GUILLET, Jean-Gerard
TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
FILE REFERENCE: 45636-5071-US
CURRENT APPLICATION NUMBER: US/10/476,570
PRIOR APPLICATION NUMBER: PCT/FR02/01533
PRIOR FILING DATE: 2002-05-03
PRIOR APPLICATION NUMBER: FR 01 05980
PRIOR FILING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 63
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 19
LENGTH: 33
TYPE: PRT
ORGANISM: artificial sequence
FEATURES:
OTHER INFORMATION: Description of the artificial sequence: peptide E6 14-46
US-10-476-570-19
```

```
Query Match          100.0%; Score 52; DB 4; Length 33;
Best Local Similarity 100.0%; Pred. No. 0.17;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 24 CVYCKQQL 32
```

```
RESULT 7
US-10-177-390-6
Sequence 6, Application US/10177390
Publication No. US20030143743A1
GENERAL INFORMATION:
APPLICANT: Schuler, Gerold
APPLICANT: N.V. Antwerp Innovatocentrum
TITLE OF INVENTION: Improved transfection of Eucaryotic Cells with linear
FILE REFERENCE: 021505wo/JH/ml
CURRENT APPLICATION NUMBER: US/10/177,390
PRIOR FILING DATE: 2002-06-20
NUMBER OF SEQ ID NOS: 34
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 6
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-177-390-6
```

```
Query Match          100.0%; Score 52; DB 4; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 30 CVYCKQQL 38
```

```
RESULT 8
US-10-484-063-20
Sequence 20, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
```

```
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLEMO
APPLICANT: FOLLEN, MICHELE
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
PRIOR FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 20
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus
US-10-484-063-20
```

```
Query Match          100.0%; Score 52; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKOQL 9
DB 30 CVYCKOQL 38
```

```
RESULT 9
US-10-484-063-27
Sequence 27, Application US/10484063
Publication No. US20050048467A1
GENERAL INFORMATION:
APPLICANT: SASTRY, K. JAGANNADHA
APPLICANT: TORTOLERO-LUNA, GUILLEMO
APPLICANT: FOLLEN, MICHELE
TITLE OF INVENTION: METHODS AND COMPOSITIONS RELATING TO HPV-ASSOCIATED
FILE REFERENCE: UTSC:560US
CURRENT APPLICATION NUMBER: US/10/484,063
CURRENT FILING DATE: 2004-01-16
PRIOR APPLICATION NUMBER: PCT/US02/23198
PRIOR FILING DATE: 2002-07-19
PRIOR APPLICATION NUMBER: 60/306,809
PRIOR FILING DATE: 2001-07-20
NUMBER OF SEQ ID NOS: 27
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 27
LENGTH: 151
TYPE: PRT
ORGANISM: Human papillomavirus type 16
US-10-484-063-27
```

```
Query Match          100.0%; Score 52; DB 5; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.66;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKOQL 9
DB 30 CVYCKOQL 38
```

```
RESULT 10
US-10-858-384-2
Sequence 2, Application US/10858384
Publication No. US2005003025A1
GENERAL INFORMATION:
APPLICANT: CHOPIIN, JEANNINE
APPLICANT: BOURGAULT VILLARD, ISABELLE
APPLICANT: GUILLET, JEAN-GERARD
APPLICANT: CONNAN, FRANCINE
APPLICANT: FERRIES, ESTELLE
```

```
TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
FILE REFERENCE: 0508-1037-1
CURRENT APPLICATION NUMBER: US/10/858,384
CURRENT FILING DATE: 2004-06-02
PRIOR APPLICATION NUMBER: FR 9907012
PRIOR FILING DATE: 1999-06-03
NUMBER OF SEQ ID NOS: 24
SOFTWARE: PatentIn Ver. 3.2
SEQ ID NO 2
LENGTH: 158
TYPE: PRT
ORGANISM: Human Papillomavirus
US-10-858-384-2
```

```
Query Match          100.0%; Score 52; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.69;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKOQL 9
DB 37 CVYCKOQL 45
```

```
RESULT 11
US-10-367-057-16
Sequence 16, Application US/10367057
Publication No. US20050100554A1
GENERAL INFORMATION:
APPLICANT: Cuthill, Scott;
APPLICANT: Jackson, Amanda;
APPLICANT: Lewin, David A.;
APPLICANT: Ooi, Chean Eng
TITLE OF INVENTION: Complexes and Methods of Using Same
FILE REFERENCE: 21402-559
CURRENT APPLICATION NUMBER: US/10/367,057
CURRENT FILING DATE: 2003-02-14
PRIOR APPLICATION NUMBER: 60/256,911
PRIOR FILING DATE: 2002-02-14
NUMBER OF SEQ ID NOS: 198
SOFTWARE: CuraSeqIst version 0.1
SEQ ID NO 16
LENGTH: 158
TYPE: PRT
ORGANISM: Homo sapiens
US-10-367-057-16
```

```
Query Match          100.0%; Score 52; DB 5; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.69;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKOQL 9
DB 37 CVYCKOQL 45
```

```
RESULT 12
US-11-021-949-13
Sequence 13, Application US/11021949
Publication No. US20050142541A1
GENERAL INFORMATION:
APPLICANT: LU, PETER
APPLICANT: GARMAN, JONATHAN DAVID
APPLICANT: BELMARES, MICHAEL P.
APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
APPLICANT: SCHWEIZER, JOHANNES
TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
FILE REFERENCE: VITA-012
CURRENT APPLICATION NUMBER: US/11/021,949
CURRENT FILING DATE: 2004-12-23
PRIOR APPLICATION NUMBER: 60/532,373
```

```
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 13
; LENGTH: 158
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-13
```

```
Query Match          100.0%; Score 52; DB 6; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.69;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
    |||||
Db 37 CVYCKQQL 45
```

```
RESULT 13
US-10-472-724-2
; Sequence 2, Application US/10472724
; Publication No. US20040171806A1
; GENERAL INFORMATION:
; APPLICANT: Cid-Arregui, Angel
; APPLICANT: Zur Hausen, Harald
; TITLE OF INVENTION: Modified HPV E6 and E7 genes and proteins useful for vaccination
; FILE REFERENCE: 4121-154
; CURRENT APPLICATION NUMBER: US/10/472,724
; PRIOR FILING DATE: 2003-09-17
; PRIOR APPLICATION NUMBER: PCT/EP02/03271
; PRIOR FILING DATE: 2002-03-22
; PRIOR APPLICATION NUMBER: EP 01107271.7
; PRIOR FILING DATE: 2001-03-23
; NUMBER OF SEQ ID NOS: 27
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 2
; LENGTH: 171
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Synthetic Construct
US-10-472-724-2
```

```
Query Match          100.0%; Score 52; DB 4; Length 171;
Best Local Similarity 100.0%; Pred. No. 0.74;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
    |||||
Db 42 CVYCKQQL 50
```

```
RESULT 14
US-11-072-288-1
; Sequence 1, Application US/11072288
; Publication No. US20050159386A1
; GENERAL INFORMATION:
; APPLICANT: KIENY, Marie-Paule
; APPLICANT: BIZOUANE, Nadine
; APPLICANT: BIZOUANE, Nadine
; TITLE OF INVENTION: ANTI-TUMORAL COMPOSITION BASED ON IMMUNOGENIC
; TITLE OF INVENTION: POLYPEPTIDE WITH MODIFIED CELL LOCATION
; FILE REFERENCE: 017753-122
; CURRENT APPLICATION NUMBER: US/11/072,288
; PRIOR FILING DATE: 2005-03-07
; PRIOR APPLICATION NUMBER: US/09/462,993
; PRIOR FILING DATE: 2000-04-17
; PRIOR APPLICATION NUMBER: PCT/FR98/01576
; PRIOR FILING DATE: 1998-07-17
; PRIOR APPLICATION NUMBER: FR 97/09152
; PRIOR FILING DATE: 1997-07-18
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: PatentIn Ver. 2.2
```

```
; SEQ ID NO 1
; LENGTH: 243
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Derived from
; OTHER INFORMATION: human papillomavirus, strain HPV-16, E6 protein
; OTHER INFORMATION: fused F protein signals, clone E6*YWF.
US-11-072-288-1
```

```
Query Match          100.0%; Score 52; DB 6; Length 243;
Best Local Similarity 100.0%; Pred. No. 1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
    |||||
Db 65 CVYCKQQL 73
```

```
RESULT 15
US-09-367-309A-1
; Sequence 1, Application US/09367309A
; Publication No. US20020081329A1
; GENERAL INFORMATION:
; APPLICANT: MACFARLAN, RODERICK I.
; APPLICANT: MALIAROS, JIM
; TITLE OF INVENTION: CHEATING IMMUNOSTIMULATING COMPLEXES
; FILE REFERENCE: 017227/0149
; CURRENT APPLICATION NUMBER: US/09/367,309A
; PRIOR FILING DATE: 1999-08-11
; PRIOR APPLICATION NUMBER: PCT/AU98/00080
; PRIOR FILING DATE: 1998-02-13
; PRIOR APPLICATION NUMBER: AU PO 5178
; PRIOR FILING DATE: 1997-02-19
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 1
; LENGTH: 266
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-09-367-309A-1
```

```
Query Match          100.0%; Score 52; DB 3; Length 266;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
    |||||
Db 37 CVYCKQQL 45
```

```
RESULT 16
US-10-000-903-4
; Sequence 4, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Gabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; PRIOR FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
```

```
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-4

Query Match          100.0%; Score 52; DB 4; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQOQL 9
Db 143 CVYCKQOQL 151

RESULT 17
US-10-899-771-4
; Sequence 4, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 4
; LENGTH: 273
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (protein D from Haemophilus
; OTHER INFORMATION: influenzae B and B6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-4

Query Match          100.0%; Score 52; DB 5; Length 273;
Best Local Similarity 100.0%; Pred. No. 1.1;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
Db 143 CVYCKOQL 151

RESULT 18
US-10-000-903-10
; Sequence 10, Application US/10000903
; Publication No. US2002018222A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
```

```
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-10

Query Match          100.0%; Score 52; DB 4; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
Db 162 CVYCKOQL 170

RESULT 19
US-10-899-771-10
; Sequence 10, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 10
; LENGTH: 292
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeric protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6 from Human papilloma virus type
; OTHER INFORMATION: 16)
US-10-899-771-10

Query Match          100.0%; Score 52; DB 5; Length 292;
Best Local Similarity 100.0%; Pred. No. 1.2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
Db 162 CVYCKOQL 170

RESULT 20
US-10-000-903-6
; Sequence 6, Application US/10000903
; Publication No. US2002018222A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabezon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Bernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 9717953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
```

```
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-6
```

```
Query Match          100.0%; Score 52; DB 4; Length 371;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVYCKQQL 9
Db 143 CVYCKQQL 151
```

```
RESULT 21
US-10-899-771-6
; Sequence 6, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 6
; LENGTH: 371
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeraic protein (protein D from Haemophilus
; OTHER INFORMATION: influenza B and B6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-6
```

```
Query Match          100.0%; Score 52; DB 5; Length 371;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVYCKQQL 9
Db 143 CVYCKQQL 151
```

```
RESULT 22
US-10-000-903-14
; Sequence 14, Application US/10000903
; Publication No. US20020182221A1
; GENERAL INFORMATION:
; APPLICANT: Bruck, Claudine
; APPLICANT: Cabazon Silva, Teresa
; APPLICANT: Delisse, Anne-Marie Eva Fernande
; APPLICANT: Gerard, Catherine Marie Ghislaine
; APPLICANT: Lombardo-Bencheikh, Angela
; TITLE OF INVENTION: Vaccine
; FILE REFERENCE: B45107
; CURRENT APPLICATION NUMBER: US/10/000,903
; CURRENT FILING DATE: 2001-10-01
; PRIOR APPLICATION NUMBER: PCT/EP98/05285
; PRIOR FILING DATE: 1998-08-17
; PRIOR APPLICATION NUMBER: GB 971953.5
; PRIOR FILING DATE: 1997-08-22
; NUMBER OF SEQ ID NOS: 23
```

```
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Homo sapien
US-10-000-903-14
```

```
Query Match          100.0%; Score 52; DB 4; Length 390;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVYCKQQL 9
Db 162 CVYCKQQL 170
```

```
RESULT 23
US-10-899-771-14
; Sequence 14, Application US/10899771
; Publication No. US20050031638A1
; GENERAL INFORMATION:
; APPLICANT: Dalemans, Wilfried L.J.
; APPLICANT: Gerard, Catherine Marie Ghislaine
; TITLE OF INVENTION: Compositions Comprising Human Papilloma Virus Proteins
; TITLE OF INVENTION: and Fusion Proteins Adjuvanted with a Cpg Oligonucleotide
; FILE REFERENCE: B45124
; CURRENT APPLICATION NUMBER: US/10/899,771
; CURRENT FILING DATE: 2004-07-27
; PRIOR APPLICATION NUMBER: US/09/581,976
; PRIOR FILING DATE: 2000-06-20
; PRIOR APPLICATION NUMBER: PCT/EP98/08563
; PRIOR FILING DATE: 1998-12-18
; PRIOR APPLICATION NUMBER: GB 9727262.9
; PRIOR FILING DATE: 1997-12-24
; NUMBER OF SEQ ID NOS: 28
; SOFTWARE: FastSeq for Windows Version 3.0
; SEQ ID NO 14
; LENGTH: 390
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Chimeraic protein (Clyta from Streptococcus
; OTHER INFORMATION: pneumoniae and E6E7 fusion from Human papilloma
; OTHER INFORMATION: virus type 16)
US-10-899-771-14
```

```
Query Match          100.0%; Score 52; DB 5; Length 390;
Best Local Similarity 100.0%; Pred. No. 1.5;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVYCKQQL 9
Db 162 CVYCKQQL 170
```

```
RESULT 24
US-10-367-095-10
; Sequence 10, Application US/10367095
; Publication No. US20030226696A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: No. US20030226696A1 Insect Cell Line
; FILE REFERENCE: 44149-1US1
; CURRENT APPLICATION NUMBER: US/10/367,095
; CURRENT FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
```

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; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FaastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-095-10

Query Match          100.0%; Score 52; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCKOQL 9
Db 507 CVCCKOQL 515

RESULT 25
US-10-368-046-10
; Sequence 10, Application US/10368046
; Publication No. US20040063188A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: Method for Isolation and Purification of
; TITLE OF INVENTION: Expressed Gene Products In Vitro
; FILE REFERENCE: 44149-1US1
; CURRENT APPLICATION NUMBER: US/10/368,046
; CURRENT FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FaastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-368-046-10
```

```

US-10-368-046-10
Query Match          100.0%; Score 52; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCKOQL 9
Db 507 CVCCKOQL 515

RESULT 26
US-10-367-367-10
; Sequence 10, Application US/10367367
; Publication No. US20040121465A1
; GENERAL INFORMATION:
; APPLICANT: Robin A. Robinson
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 44149-2US1
; CURRENT APPLICATION NUMBER: US/10/367,367
; CURRENT FILING DATE: 2003-02-15
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,118
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,133
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FaastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-367-367-10

Query Match          100.0%; Score 52; DB 4; Length 536;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVCCKOQL 9
Db 507 CVCCKOQL 515

RESULT 27
US-10-918-337-10
; Sequence 10, Application US/10918337
; Publication No. US20050118191A1
; GENERAL INFORMATION:
; APPLICANT: NOVAVAX, INC., et al.
; TITLE OF INVENTION: Optimization of Gene Sequences of
; TITLE OF INVENTION: Chimeric Virus-Like Particles for Expression in Insect Cells
; FILE REFERENCE: 19065/2132
; CURRENT APPLICATION NUMBER: US/10/918,337
; CURRENT FILING DATE: 2004-08-13
; PRIOR APPLICATION NUMBER: PCT/US03/04473
; PRIOR FILING DATE: 2003-02-14
; PRIOR APPLICATION NUMBER: US 60/356,119
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,161
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,135
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FaastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10
```

```

; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,157
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,156
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,123
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,113
; PRIOR FILING DATE: 2002-02-14
; PRIOR APPLICATION NUMBER: US 60/356,154
; PRIOR FILING DATE: 2002-02-14
; Remaining Prior Application data removed - See file wrapper or PALM.
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 10
; LENGTH: 536
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: HPV-16 L2/E6 fusion protein
US-10-918-337-10
```

```

Query Match          100.0%; Score 52; DB 5; Length 536;
Best Local Similarity 100.0%; Pred. No. 2;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVCCKOQL 9
Db 507 CVCCKOQL 515
```

```

RESULT 28
US-10-476-570-53
; Sequence 53, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLIERE, Bernard
; APPLICANT: BOURGAULT-VILLADA, Isabelle
; APPLICANT: POUVELLE-MORITILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; PRIOR FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 53
; LENGTH: 30
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 15-44
US-10-476-570-53
```

```

Query Match          92.3%; Score 48; DB 4; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVCCKOQL 8
Db 23 CVCCKOQL 30
```

```

RESULT 29
US-10-858-384-4
; Sequence 4, Application US/10858384
```

```

; Publication No. US2005033025A1
; GENERAL INFORMATION:
; APPLICANT: CHOPPIN, JEANNINE
; APPLICANT: BOURGAULT VILLADA, ISABELLE
; APPLICANT: GUILLET, JEAN-GERARD
; APPLICANT: CONNAN, FRANCINE
; APPLICANT: FERRIES, ESTELLE
; TITLE OF INVENTION: POLYPEPTIDIC PROTEIN FRAGMENTS OF THE E6 PROTEIN
; TITLE OF INVENTION: OR E7 OF HPV, THEIR PRODUCTION AND THEIR USE
; FILE REFERENCE: 0508-1037-1
; CURRENT APPLICATION NUMBER: US/10/858,384
; PRIOR FILING DATE: 2004-06-02
; PRIOR APPLICATION NUMBER: FR 9907012
; PRIOR FILING DATE: 1999-06-03
; NUMBER OF SEQ ID NOS: 24
; SOFTWARE: PatentIn Ver. 3.2
; SEQ ID NO 4
; LENGTH: 30
; TYPE: PRT
; ORGANISM: Artificial Sequence
; FEATURE:
; OTHER INFORMATION: Description of the Artificial Sequence: Peptide fragment
US-10-858-384-4
```

```

Query Match          92.3%; Score 48; DB 5; Length 30;
Best Local Similarity 100.0%; Pred. No. 0.72;
Matches 8; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVCCKOQL 8
Db 23 CVCCKOQL 30
```

```

RESULT 30
US-11-021-949-14
; Sequence 14, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BRIMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; PRIOR FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 14
; LENGTH: 149
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-14
```

```

Query Match          86.5%; Score 45; DB 6; Length 149;
Best Local Similarity 87.5%; Pred. No. 9.3;
Matches 7; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
```

```
Qy 1 CVCCKOQL 8
Db 30 CVCCKOQL 37
```

```

RESULT 31
US-11-021-949-361
; Sequence 361, Application US/11021949
; Publication No. US20050142541A1
```

```
/ GENERAL INFORMATION:
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
/ APPLICANT: SCHWEIZER, JOHANNES
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
/ PRIOR FILING DATE: 2004-12-23
/ PRIOR APPLICATION NUMBER: 60/532,373
/ NUMBER OF SEQ ID NOS: 361
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 361
/ LENGTH: 158
/ TYPE: PRT
/ ORGANISM: human papilloma virus (HPV)
US-11-021-949-361
```

```
Query Match      80.8%; Score 42; DB 6; Length 158;
Best Local Similarity 87.5%; Pred. No. 31;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 CVCCKOOL 8
        |||||:|
DB      32 CVCCKOOL 39
```

```
RESULT 32
US-11-021-949-18
/ Sequence 18, Application US/11021949
/ Publication No. US20050142541A1
/ GENERAL INFORMATION:
```

```
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
/ APPLICANT: SCHWEIZER, JOHANNES
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
/ PRIOR FILING DATE: 2004-12-23
/ PRIOR APPLICATION NUMBER: 60/532,373
/ NUMBER OF SEQ ID NOS: 361
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 18
/ LENGTH: 149
/ TYPE: PRT
/ ORGANISM: human papilloma virus (HPV)
US-11-021-949-18
```

```
Query Match      78.8%; Score 41; DB 6; Length 149;
Best Local Similarity 87.5%; Pred. No. 42;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 CVCCKOOL 8
        |||||:|
DB      30 CVCCKOOL 37
```

```
RESULT 33
US-11-021-949-24
/ Sequence 24, Application US/11021949
/ Publication No. US20050142541A1
/ GENERAL INFORMATION:
```

```
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
```

```
/ APPLICANT: SCHWEIZER, JOHANNES
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
/ PRIOR FILING DATE: 2004-12-23
/ PRIOR APPLICATION NUMBER: 60/532,373
/ NUMBER OF SEQ ID NOS: 361
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 24
/ LENGTH: 151
/ TYPE: PRT
/ ORGANISM: human papilloma virus (HPV)
US-11-021-949-24
```

```
Query Match      78.8%; Score 41; DB 6; Length 151;
Best Local Similarity 75.0%; Pred. No. 43;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVCCKOOL 8
        |||||:|
DB      30 CVCCKOOL 37
```

```
RESULT 34
US-11-021-949-25
/ Sequence 25, Application US/11021949
/ Publication No. US20050142541A1
/ GENERAL INFORMATION:
```

```
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
/ APPLICANT: SCHWEIZER, JOHANNES
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
/ PRIOR FILING DATE: 2004-12-23
/ PRIOR APPLICATION NUMBER: 60/532,373
/ NUMBER OF SEQ ID NOS: 361
/ SOFTWARE: FastSeq for Windows Version 4.0
/ SEQ ID NO 25
/ LENGTH: 151
/ TYPE: PRT
/ ORGANISM: human papilloma virus (HPV)
US-11-021-949-25
```

```
Query Match      78.8%; Score 41; DB 6; Length 151;
Best Local Similarity 75.0%; Pred. No. 43;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVCCKOOL 8
        |||||:|
DB      30 CVCCKOOL 37
```

```
RESULT 35
US-11-021-949-20
/ Sequence 20, Application US/11021949
/ Publication No. US20050142541A1
/ GENERAL INFORMATION:
```

```
/ APPLICANT: LU, PETER
/ APPLICANT: GARMAN, JONATHAN DAVID
/ APPLICANT: BELMARES, MICHAEL P.
/ APPLICANT: DIAZ-SARMIENTO, CHAMORRO SONOZA
/ APPLICANT: SCHWEIZER, JOHANNES
/ TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
/ FILE REFERENCE: VITA-012
/ CURRENT APPLICATION NUMBER: US/11/021,949
```



```

; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 20
; LENGTH: 153
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-20
```

```

Query Match      78.8%; Score 41; DB 6; Length 153;
Best Local Similarity 75.0%; Pred. No. 43;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1  CVYCKQOL 8
Db      33  CVYCKKEL 40
```

```

RESULT 36
US-11-021-949-22
; Sequence 22, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 22
; LENGTH: 155
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-22
```

```

Query Match      78.8%; Score 41; DB 6; Length 155;
Best Local Similarity 75.0%; Pred. No. 44;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1  CVYCKQOL 8
Db      33  CVYCKKEL 40
```

```

RESULT 37
US-11-021-949-23
; Sequence 23, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
```

```

; SEQ ID NO 23
; LENGTH: 155
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-23
```

```

Query Match      78.8%; Score 41; DB 6; Length 155;
Best Local Similarity 75.0%; Pred. No. 44;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1  CVYCKQOL 8
Db      33  CVYCKKEL 40
```

```

RESULT 38
US-11-021-949-31
; Sequence 31, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; TITLE OF INVENTION: AND METHODS OF THEIR USE
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 31
; LENGTH: 162
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-31
```

```

Query Match      78.8%; Score 41; DB 6; Length 162;
Best Local Similarity 75.0%; Pred. No. 46;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1  CVYCKQOL 8
Db      36  CVYCKRQL 43
```

```

RESULT 39
US-10-437-963-106779
; Sequence 106779, Application US/10437963
; Publication No. US20040123343A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; APPLICANT: Wu, Wei
; APPLICANT: Boukharov, Andrey A.
; APPLICANT: Barbazuk, Brad
; APPLICANT: Li, Ping
; TITLE OF INVENTION: Rice Nucleic Acid Molecules and Other Molecules Associated With
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53221)B
; CURRENT APPLICATION NUMBER: US/10/437,963
; CURRENT FILING DATE: 2003-05-14
; NUMBER OF SEQ ID NOS: 204966
; SEQ ID NO 106779
; LENGTH: 135
; TYPE: PRT
; ORGANISM: Oryza sativa
; FEATURES:
```

```

; NAME/KEY: unsure
; LOCATION: (1)..(135)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT4530_11193C.1.pep
US-10-437-963-106779
```

```
Query Match          76.9%; Score 40; DB 4; Length 135;
Best Local Similarity 85.7%; Pred. No. 57;
Matches 6; Conservative 1; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQO 7
        ||:||||
DB      30 CVYCKQO 36
```

```
RESULT 40
US-11-021-949-359
; Sequence 359, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: PaacSeq for Windows Version 4.0
; SEQ ID NO 359
; LENGTH: 148
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-359
```

```
Query Match          76.9%; Score 40; DB 6; Length 148;
Best Local Similarity 75.0%; Pred. No. 62;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 8
        |||::||
DB      31 CVYCEROL 38
```

```
RESULT 41
US-10-476-570-23
; Sequence 23, Application US/10476570
; Publication No. US20040170644A1
; GENERAL INFORMATION:
; APPLICANT: COMMISSARIAT A L'ENERGIE ATOMIQUE
; APPLICANT: INSTITUT NATIONAL DE LA SANTE ET DE LA RECHERCHE MEDICALE
; APPLICANT: MAILLERE, Bernard
; APPLICANT: BOURGAULT-VILADA, Isabelle
; APPLICANT: ROUYABLE-MORATILLE, Sandra
; APPLICANT: GUILLET, Jean-Gerard
; TITLE OF INVENTION: Mixture of peptides derived from E6 and/or E7
; FILE REFERENCE: 45636-5071-US
; CURRENT APPLICATION NUMBER: US/10/476,570
; CURRENT FILING DATE: 2003-11-04
; PRIOR APPLICATION NUMBER: PCT/FR02/01533
; PRIOR FILING DATE: 2002-05-03
; PRIOR APPLICATION NUMBER: FR 01 05980
; PRIOR FILING DATE: 2001-05-04
; NUMBER OF SEQ ID NOS: 63
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 23
```

```

; LENGTH: 15
; TYPE: PRT
; ORGANISM: artificial sequence
; FEATURE:
; OTHER INFORMATION: Description of the artificial sequence: peptide E6 28-42
US-10-476-570-23
```

```
Query Match          75.0%; Score 39; DB 4; Length 15;
Best Local Similarity 100.0%; Pred. No. 12;
Matches 6; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQ 6
        ||||||
DB      10 CVYCKQ 15
```

```
RESULT 42
US-10-369-493-11069
; Sequence 11069, Application US/10369493
; Publication No. US20030233675A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEQ ID NO 11069
; LENGTH: 147
; TYPE: PRT
; ORGANISM: Ferrioplasma acidarmanus
US-10-369-493-11069
```

```
Query Match          75.0%; Score 39; DB 4; Length 147;
Best Local Similarity 62.5%; Pred. No. 89;
Matches 5; Conservative 3; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 8
        |||::||
DB      131 CVYCKKEM 138
```

```
RESULT 43
US-11-021-949-27
; Sequence 27, Application US/11021949
; Publication No. US20050142541A1
; GENERAL INFORMATION:
; APPLICANT: LU, PETER
; APPLICANT: GARMAN, JONATHAN DAVID
; APPLICANT: BELMARES, MICHAEL P.
; APPLICANT: DIAZ-SARMIENTO, CHAMORRO SOMOZA
; APPLICANT: SCHWEIZER, JOHANNES
; TITLE OF INVENTION: ANTIBODIES FOR ONCOGENIC STRAINS OF HPV
; FILE REFERENCE: VITA-012
; CURRENT APPLICATION NUMBER: US/11/021,949
; CURRENT FILING DATE: 2004-12-23
; PRIOR APPLICATION NUMBER: 60/532,373
; PRIOR FILING DATE: 2003-12-23
; NUMBER OF SEQ ID NOS: 361
; SOFTWARE: PaacSeq for Windows Version 4.0
; SEQ ID NO 27
; LENGTH: 150
; TYPE: PRT
; ORGANISM: human papilloma virus (HPV)
US-11-021-949-27
```

Query Match 75.0%; Score 39; DB 6; Length 150;
Best Local Similarity 75.0%; Pred. No. 91;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQOL 8
DB 30 CVYCKETL 37

RESULT 44
US-10-424-599-229548
; Sequence 229548, Application US/10424599
; Publication No. US20040031072A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa Thomas J
; APPLICANT: Kovalic David K
; APPLICANT: Zhou Yihua
; APPLICANT: Cao Yongwei
; TITLE OF INVENTION: Soy Nucleic Acid Molecules and Other Molecules Associated with
; TITLE OF INVENTION: Plants and Uses Thereof for Plant Improvement
; FILE REFERENCE: 38-21(53223)B
; CURRENT APPLICATION NUMBER: US/10/424,599
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 285684
; SEQ ID NO 229548
; LENGTH: 157
; TYPE: PRT
; ORGANISM: Glycine max
; FEATURE:
; OTHER INFORMATION: Clone ID: PAT_MRT3847_49306C.1.pep
US-10-424-599-229548

Query Match 75.0%; Score 39; DB 4; Length 157;
Best Local Similarity 75.0%; Pred. No. 95;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQOL 8
DB 93 CVYCKQKL 100

RESULT 45
US-10-425-115-312020
; Sequence 312020, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovalic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated with
; TITLE OF INVENTION: Plants
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326
; SEQ ID NO 312020
; LENGTH: 185
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(185)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_47614C.1.pep
US-10-425-115-312020

Query Match 75.0%; Score 39; DB 4; Length 185;
Best Local Similarity 75.0%; Pred. No. 11e+02;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQOL 8
DB 133 CVYCKQKL 140

RESULT 46
US-10-291-737-6
; Sequence 6, Application US/10291737
; Publication No. US20030087299A1
; GENERAL INFORMATION:
; APPLICANT: WEBSTER, Marion et al.
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: AND USES THEREOF
; FILE REFERENCE: CL001017-CON
; CURRENT APPLICATION NUMBER: US/10/291,737
; CURRENT FILING DATE: 2002-11-12
; PRIOR APPLICATION NUMBER: US 09/822,859
; PRIOR FILING DATE: 2001-04-02
; PRIOR APPLICATION NUMBER: US 60/254,588
; PRIOR FILING DATE: 2000-12-12
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6
; LENGTH: 398
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-10-291-737-6

Query Match 75.0%; Score 39; DB 4; Length 398;
Best Local Similarity 55.6%; Pred. No. 2.2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQOL 9
DB 41 CIVCTROLI 49

RESULT 47
US-10-365-564-6
; Sequence 6, Application US/10365564
; Publication No. US20030143623A1
; GENERAL INFORMATION:
; APPLICANT: WEBSTER, Marion et al.
; TITLE OF INVENTION: ISOLATED HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: NUCLEIC ACID MOLECULES ENCODING HUMAN TRANSPORTER PROTEINS,
; TITLE OF INVENTION: AND USES THEREOF
; FILE REFERENCE: CL001017-CIPCON
; CURRENT APPLICATION NUMBER: US/10/365,564
; CURRENT FILING DATE: 2003-02-13
; PRIOR APPLICATION NUMBER: US 09/822,859
; PRIOR FILING DATE: 2001-04-02
; PRIOR APPLICATION NUMBER: US 60/254,588
; PRIOR FILING DATE: 2000-12-12
; NUMBER OF SEQ ID NOS: 6
; SOFTWARE: FastSeq for Windows Version 4.0
; SEQ ID NO 6
; LENGTH: 398
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-10-365-564-6

Query Match 75.0%; Score 39; DB 4; Length 398;
Best Local Similarity 55.6%; Pred. No. 2.2e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

QY 1 CVYCKQOL 9
DB 41 CIVCTROLI 49

RESULT 48
US-10-225-810-35

```

; Sequence 35, Application US/10225810
; Publication No. US20030157512A1
; GENERAL INFORMATION:
; APPLICANT: Bermingham, Jr., John R.
; TITLE OF INVENTION: Transdorins and Methods of Using Transdorin
; FILE REFERENCE: McLaugh-07165
; CURRENT APPLICATION NUMBER: US/10/225,810
; NUMBER OF SEQ ID NOS: 76
; SOFTWARE: PatentIn version 3.2
; SEQ ID NO 35
; LENGTH: 449
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-10-225-810-35

```

```

Query Match          75.0%; Score 39; DB 4; Length 449;
Best Local Similarity 55.6%; Pred. No. 2.4e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 CVYCKQQL 9
|:|:|:|:
DB      83 CICYCTRL 91

```

```

RESULT 49
US-10-369-493-5326
; Sequence 5326, Application US/10369493
; Publication No. US20030233675A1
; GENERAL INFORMATION:
; APPLICANT: Cao, Yongwei
; APPLICANT: Hinkle, Gregory J.
; APPLICANT: Slater, Steven C.
; APPLICANT: Goldman, Barry S.
; APPLICANT: Chen, Xianfeng
; TITLE OF INVENTION: EXPRESSION OF MICROBIAL PROTEINS IN PLANTS FOR PRODUCTION OF
; FILE REFERENCE: 38-10(52052)B
; CURRENT APPLICATION NUMBER: US/10/369,493
; CURRENT FILING DATE: 2003-02-28
; PRIOR APPLICATION NUMBER: US 60/360,039
; PRIOR FILING DATE: 2002-02-21
; NUMBER OF SEQ ID NOS: 47374
; SEQ ID NO 5326
; LENGTH: 449
; TYPE: PRT
; ORGANISM: Caenorhabditis elegans
US-10-369-493-5326

```

```

Query Match          75.0%; Score 39; DB 4; Length 449;
Best Local Similarity 55.6%; Pred. No. 2.4e+02;
Matches 5; Conservative 3; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      1 CVYCKQQL 9
|:|:|:|:
DB      83 CICYCTRL 91

```

```

RESULT 50
US-10-425-115-199640
; Sequence 199640, Application US/10425115
; Publication No. US20040214272A1
; GENERAL INFORMATION:
; APPLICANT: La Rosa, Thomas J.
; APPLICANT: Kovacic, David K.
; APPLICANT: Zhou, Yihua
; APPLICANT: Cao, Yongwei
; TITLE OF INVENTION: Nucleic Acid Molecules and Other Molecules Associated With
; FILE REFERENCE: 38-21(53222)B
; CURRENT APPLICATION NUMBER: US/10/425,115
; CURRENT FILING DATE: 2003-04-28
; NUMBER OF SEQ ID NOS: 369326

```

```

; SEQ ID NO 199640
; LENGTH: 145
; TYPE: PRT
; ORGANISM: Zea mays
; FEATURE:
; NAME/KEY: unsure
; LOCATION: (1)..(145)
; OTHER INFORMATION: unsure at all Xaa locations
; FEATURE:
; OTHER INFORMATION: Clone ID: MRT4577_113644C.1.pep
US-10-425-115-199640

```

```

Query Match          73.1%; Score 38; DB 4; Length 145;
Best Local Similarity 87.5%; Pred. No. 1.3e+02;
Matches 7; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

```

```

QY      2 VYCKQQL 9
|:|:|:|:
DB      63 VYCKQQL 70

```

```

Search completed: May 5, 2006, 08:28:54
Job time : 58 secs

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GenCore version 5.1.7
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OM protein - protein search, using SW model

Run on: May 5, 2006, 08:19:00 ; Search time 8.4 Seconds
(without alignments)
49.591 Million cell updates/sec

Title: US-08-170-344-9
Perfect score: 52
Sequence: 1 CVYCKQQLL 9

Scoring table: BLOSUM62
Gapop 10.0 , Gapext 0.5

Searched: 235405 seqs, 46284737 residues

Total number of hits satisfying chosen parameters: 235405

Minimum DB seq length: 0
Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 1000 summaries

Database :

Published Applications AA New:*

- 1: /SIDS5/pcodata/1/pubppaa/US08_NEW_PUB.pep1.*
- 2: /SIDS5/pcodata/1/pubppaa/US06_NEW_PUB.pep.*
- 3: /SIDS5/pcodata/1/pubppaa/US07_NEW_PUB.pep.*
- 4: /SIDS5/pcodata/1/pubppaa/US08_NEW_PUB.pep.*
- 5: /SIDS5/pcodata/1/pubppaa/PCCT_NEW_PUB.pep.*
- 6: /SIDS5/pcodata/1/pubppaa/US09_NEW_PUB.pep.*
- 7: /SIDS5/pcodata/1/pubppaa/US09_NEW_PUB.pep1.*
- 8: /SIDS5/pcodata/1/pubppaa/US10_NEW_PUB.pep.*
- 9: /SIDS5/pcodata/1/pubppaa/US10_NEW_PUB.pep1.*
- 10: /SIDS5/pcodata/1/pubppaa/US11_NEW_PUB.pep1.*
- 11: /SIDS5/pcodata/1/pubppaa/US11_NEW_PUB.pep1.*
- 12: /SIDS5/pcodata/1/pubppaa/US60_NEW_PUB.pep.*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	Query Match	Length	DB ID	Description
1	52	100.0	151	9	US-10-530-253-13
2	52	100.0	158	11	US-11-206-138-3
3	52	100.0	248	9	US-10-530-253-1
4	52	100.0	248	9	US-10-530-253-3
5	52	100.0	248	9	US-10-530-253-5
6	52	100.0	248	9	US-10-530-253-7
7	52	100.0	248	9	US-10-530-253-9
8	52	100.0	248	9	US-10-530-253-11
9	52	100.0	256	11	US-11-192-923A-2
10	45	86.5	149	9	US-10-530-253-18
11	41	78.8	15	9	US-10-530-061-1692
12	41	78.8	149	9	US-10-530-253-16
13	41	78.8	151	9	US-10-530-253-21
14	41	78.8	155	9	US-10-530-253-23
15	41	78.8	158	9	US-10-530-253-26
16	40	76.9	152	9	US-10-530-253-39
17	38	73.1	351	11	US-11-087-099-8245
18	38	73.1	351	11	US-11-087-099-8245
19	37	71.2	269	9	US-10-453-372-36
20	36	69.2	10	9	US-10-530-061-562
21	36	69.2	158	9	US-10-530-253-20

22	36	69.2	343	11	US-11-087-099-7737	Sequence 7737, Ap
23	36	69.2	456	11	US-11-087-099-3798	Sequence 3798, Ap
24	36	69.2	456	11	US-11-087-099-6259	Sequence 6259, Ap
25	36	69.2	456	11	US-11-087-099-9642	Sequence 9642, Ap
26	36	69.2	456	11	US-11-188-298-14550	Sequence 14550, A
27	36	69.2	456	11	US-11-188-298-14958	Sequence 14958, A
28	36	69.2	457	11	US-11-087-099-5321	Sequence 5321, Ap
29	36	69.2	457	11	US-11-087-099-5761	Sequence 5761, Ap
30	36	69.2	457	11	US-11-087-099-9118	Sequence 9118, Ap
31	36	69.2	457	11	US-11-087-099-10468	Sequence 10468, A
32	36	69.2	457	11	US-11-087-099-11070	Sequence 11070, A
33	36	69.2	457	11	US-11-188-298-4860	Sequence 4860, Ap
34	36	69.2	457	11	US-11-188-298-8450	Sequence 8450, Ap
35	36	69.2	457	11	US-11-188-298-10221	Sequence 10221, A
36	36	69.2	457	11	US-11-188-298-16296	Sequence 16296, A
37	36	69.2	457	11	US-11-188-298-20671	Sequence 20671, A
38	36	69.2	465	11	US-11-087-099-1964	Sequence 1964, Ap
39	36	69.2	465	11	US-11-188-298-1940	Sequence 1940, Ap
40	36	69.2	466	11	US-11-087-099-8259	Sequence 8259, Ap
41	36	69.2	466	11	US-11-188-298-18632	Sequence 18632, A
42	35	67.3	9	9	US-10-530-061-621	Sequence 621, App
43	35	67.3	11	9	US-10-530-061-493	Sequence 493, App
44	35	67.3	35	11	US-11-004-399-2142	Sequence 2142, Ap
45	35	67.3	109	11	US-11-188-298-13134	Sequence 13134, A
46	35	67.3	158	9	US-10-530-253-15	Sequence 15, App1
47	35	67.3	158	9	US-10-530-253-19	Sequence 19, App1
48	35	67.3	343	11	US-11-087-099-5274	Sequence 5274, Ap
49	35	67.3	343	11	US-11-045-004-2215	Sequence 2215, Ap
50	34	65.4	10	9	US-10-530-061-560	Sequence 560, App
51	34	65.4	15	9	US-10-530-061-1659	Sequence 1659, Ap
52	34	65.4	15	9	US-10-530-061-1691	Sequence 1691, Ap
53	34	65.4	122	11	US-11-079-663-5531	Sequence 5531, Ap
54	34	65.4	160	11	US-11-096-568A-9698	Sequence 9698, Ap
55	34	65.4	160	9	US-10-530-253-25	Sequence 25, App1
56	34	65.4	176	11	US-11-096-568A-9697	Sequence 9697, Ap
57	34	65.4	320	9	US-10-453-372-854	Sequence 854, App
58	34	65.4	844	9	US-10-453-372-852	Sequence 852, App
59	34	65.4	844	9	US-10-453-372-856	Sequence 856, App
60	33	63.5	14	9	US-10-895-064-284	Sequence 284, App
61	33	63.5	14	11	US-11-116-144-212	Sequence 212, App
62	33	63.5	14	11	US-11-116-144-214	Sequence 214, App
63	33	63.5	14	11	US-11-116-144-215	Sequence 215, App
64	33	63.5	14	11	US-11-116-144-243	Sequence 243, App
65	33	63.5	14	11	US-11-129-741-284	Sequence 284, App
66	33	63.5	14	11	US-11-129-741-3216	Sequence 3216, App
67	33	63.5	14	11	US-11-220-372-212	Sequence 212, App
68	33	63.5	14	11	US-11-220-372-214	Sequence 214, App
69	33	63.5	14	11	US-11-220-372-215	Sequence 215, App
70	33	63.5	14	11	US-11-220-372-243	Sequence 243, App
71	33	63.5	28	9	US-10-729-121-17	Sequence 17, App1
72	33	63.5	28	11	US-11-285-537-17	Sequence 17, App1
73	33	63.5	37	11	US-10-895-064-2682	Sequence 2682, Ap
74	33	63.5	37	11	US-11-129-741-2682	Sequence 2682, Ap
75	33	63.5	148	9	US-10-506-454-778	Sequence 778, App
76	33	63.5	358	9	US-10-506-454-307	Sequence 307, App
77	33	63.5	374	9	US-10-506-454-908	Sequence 908, App
78	33	63.5	389	11	US-11-096-568A-15113	Sequence 15113, A
79	33	63.5	408	11	US-11-096-568A-15112	Sequence 15112, A
80	33	63.5	444	11	US-11-072-512-3354	Sequence 2354, Ap
81	33	63.5	444	11	US-11-188-298-21079	Sequence 21079, A
82	33	63.5	479	11	US-11-096-568A-15111	Sequence 15111, A
83	33	63.5	482	11	US-11-129-442-47	Sequence 47, App1
84	33	63.5	537	11	US-11-129-442-47	Sequence 16, App1
85	33	63.5	567	11	US-11-129-442-47	Sequence 9603, Ap
86	33	63.5	585	11	US-11-079-463-8980	Sequence 8980, Ap
87	33	63.5	140	11	US-11-079-463-8980	Sequence 9603, Ap
88	32	61.5	177	11	US-11-096-568A-19788	Sequence 19788, A
89	32	61.5	178	11	US-11-096-568A-3956	Sequence 3956, Ap
90	32	61.5	179	11	US-11-096-568A-34204	Sequence 34204, A
91	32	61.5	214	11	US-11-096-568A-19787	Sequence 19787, A
92	32	61.5	220	11	US-11-096-568A-34203	Sequence 34203, A
93	32	61.5	248	11	US-11-096-568A-3955	Sequence 3955, A
94	32	61.5	248	11	US-11-096-568A-3955	Sequence 3955, A

95	32	61.5	248	11	US-11-096-568A-19786	Sequence 19786, A	168	31	59.6	386	8	US-10-511-937-2930	Sequence 2930, Ap
96	32	61.5	249	11	US-11-096-568A-34202	Sequence 34202, A	169	31	59.6	415	11	US-11-182-946-6	Sequence 6, Appl1
97	32	61.5	278	9	US-10-506-454-935	Sequence 935, App	170	31	59.6	432	11	US-11-264-096-2238	Sequence 2238, Ap
98	32	61.5	309	11	US-11-087-099-2169	Sequence 2169, Ap	171	31	59.6	432	11	US-11-264-096-2239	Sequence 2239, Ap
99	32	61.5	349	11	US-11-087-099-1673	Sequence 1673, Ap	172	31	59.6	438	9	US-10-641-678-49	Sequence 49, Appl1
100	32	61.5	349	11	US-11-087-099-3545	Sequence 3545, Ap	173	31	59.6	438	11	US-11-074-176-152	Sequence 152, App
101	32	61.5	349	11	US-11-087-099-7966	Sequence 7966, Ap	174	31	59.6	438	11	US-11-079-463-9043	Sequence 9043, Ap
102	32	61.5	354	11	US-11-188-298-16534	Sequence 16534, A	175	31	59.6	456	11	US-11-188-298-12690	Sequence 12690, A
103	32	61.5	358	11	US-11-098-686-11346	Sequence 11346, A	176	31	59.6	567	11	US-11-188-298-4549	Sequence 4549, Ap
104	32	61.5	416	11	US-11-195-851-18	Sequence 18, Appl1	177	31	59.6	578	9	US-10-763-712A-4	Sequence 34, Appl1
105	32	61.5	417	9	US-10-194-487-474	Sequence 474, App	178	31	59.6	769	11	US-11-188-298-18431	Sequence 18431, A
106	32	61.5	417	9	US-10-195-883-474	Sequence 474, App	179	31	59.6	795	11	US-11-072-512-2378	Sequence 2378, Ap
107	32	61.5	417	9	US-10-195-888-474	Sequence 474, App	180	31	59.6	808	11	US-11-072-512-3401	Sequence 3401, Ap
108	32	61.5	417	9	US-10-195-889-474	Sequence 474, App	181	31	59.6	819	11	US-11-124-367A-329	Sequence 329, App
109	32	61.5	417	9	US-10-218-784-220	Sequence 220, App	182	31	59.6	912	11	US-11-124-367A-328	Sequence 328, App
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121	31	59.6	10	9	US-10-530-061-517	Sequence 517, App	194	30	57.7	105	11	US-11-264-096-232	Sequence 232, App
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127	31	59.6	142	11	US-11-225-709-2	Sequence 2, Appl1	200	30	57.7	119	11	US-11-106-399-10	Sequence 10, Appl1
128	31	59.6	143	11	US-11-196-618-23	Sequence 23, Appl1	201	30	57.7	197	11	US-11-079-463-33678	Sequence 33678, A
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130	31	59.6	178	11	US-11-096-568A-11957	Sequence 11957, A	203	30	57.7	218	11	US-11-180-990-22	Sequence 2, Appl1
131	31	59.6	194	9	US-10-793-626-140	Sequence 140, App	204	30	57.7	237	9	US-10-519-397-82	Sequence 22, Appl1
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137	31	59.6	281	11	US-11-188-298-17151	Sequence 17151, A	210	30	57.7	227	11	US-11-176-830-765	Sequence 765, App
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152	31	59.6	324	11	US-11-196-618-16	Sequence 16, Appl1	225	30	57.7	227	11	US-11-176-830-780	Sequence 780, App
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155	31	59.6	333	8	US-10-511-455-26	Sequence 26, Appl1	228	30	57.7	227	11	US-11-176-830-783	Sequence 783, App
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255	30	57.7	352	11	US-11-087-099-10074	Sequence 10074, A	328	29	55.8	416	9	US-11-072-512-3885	Sequence 3885, Ap
256	30	57.7	353	11	US-11-188-298-4314	Sequence 4314, Ap	329	29	55.8	420	11	US-11-096-568A-21736	Sequence 21736, Ap
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262	30	57.7	452	8	US-10-505-928-621	Sequence 621, App	335	29	55.8	481	11	US-11-227-177-1	Sequence 1, App1
263	30	57.7	484	11	US-11-096-568A-7428	Sequence 7428, Ap	336	29	55.8	505	11	US-11-264-096-808	Sequence 808, App
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269	30	57.7	1446	11	US-11-188-298-9230	Sequence 9230, Ap	342	29	55.8	602	9	US-10-467-657-1782	Sequence 1782, Ap
270	29	55.8	49	11	US-11-229-769-222	Sequence 222, App	343	29	55.8	604	11	US-11-183-136-4	Sequence 4, App1
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279	29	55.8	117	11	US-11-037-199-23	Sequence 23, App1	352	29	55.8	787	11	US-11-087-099-3299	Sequence 3299, Ap
280	29	55.8	135	9	US-10-523-362-32	Sequence 32, App1	353	29	55.8	787	11	US-11-188-298-633	Sequence 633, App
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297	29	55.8	232	11	US-11-096-568A-16025	Sequence 16025, A	370	28	53.8	107	11	US-11-072-512-2122	Sequence 2122, App
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394	28	53.8	262	11	US-11-072-512-2104	Sequence 2104, App	467	28	53.8	504	11	US-11-096-568a-21813	Sequence 21813, A
395	28	53.8	263	11	US-11-098-686-10590	Sequence 10590, A	468	28	53.8	504	11	US-11-079-463-7855	Sequence 7855, App
396	28	53.8	269	11	US-11-096-568a-32810	Sequence 32810, A	469	28	53.8	510	11	US-11-079-463-9001	Sequence 9001, App
397	28	53.8	271	9	US-10-533-811-1	Sequence 1, App1	470	28	53.8	511	11	US-11-096-568a-30209	Sequence 30209, A
398	28	53.8	279	8	US-10-505-928-623	Sequence 623, App	471	28	53.8	512	11	US-11-096-568a-30208	Sequence 30208, A
399	28	53.8	284	11	US-11-124-367a-382	Sequence 382, App	472	28	53.8	513	9	US-10-063-703-124	Sequence 536, App
400	28	53.8	286	11	US-11-098-686-10566	Sequence 10566, A	473	28	53.8	513	9	US-10-973-115B-536	Sequence 536, App
401	28	53.8	296	11	US-11-079-463-6624	Sequence 6624, App	474	28	53.8	513	9	US-10-137-873a-536	Sequence 536, App
402	28	53.8	297	11	US-11-079-463-8884	Sequence 8884, App	475	28	53.8	513	9	US-10-137-873a-536	Sequence 536, App
403	28	53.8	299	9	US-10-506-454-1001	Sequence 1001, App	476	28	53.8	513	9	US-10-152-370-536	Sequence 124, App
404	28	53.8	306	11	US-11-087-099-11735	Sequence 11735, A	477	28	53.8	513	11	US-11-102-240-124	Sequence 124, App
405	28	53.8	309	11	US-11-087-099-7821	Sequence 7821, App	478	28	53.8	513	11	US-11-103-195-124	Sequence 536, App
406	28	53.8	310	11	US-11-010-239-97	Sequence 97, App1	479	28	53.8	513	11	US-11-290-153-536	Sequence 536, App
407	28	53.8	310	11	US-11-087-099-11476	Sequence 11476, A	480	28	53.8	521	11	US-11-079-463-8817	Sequence 8817, App
408	28	53.8	319	11	US-11-045-004-2130	Sequence 2130, App	481	28	53.8	528	11	US-11-188-298-2430	Sequence 2430, App
409	28	53.8	326	11	US-11-087-099-10002	Sequence 10002, A	482	28	53.8	533	9	US-10-330-773-666	Sequence 666, App
410	28	53.8	338	11	US-11-087-099-276	Sequence 276, App	483	28	53.8	537	9	US-10-330-773-263	Sequence 263, App
411	28	53.8	333	11	US-11-087-099-4294	Sequence 4294, App	484	28	53.8	548	11	US-11-188-298-5263	Sequence 5263, App
412	28	53.8	339	11	US-11-188-298-2603	Sequence 2603, App	485	28	53.8	566	11	US-11-079-463-7770	Sequence 7770, App
413	28	53.8	341	11	US-11-087-099-10839	Sequence 10839, A	486	28	53.8	567	11	US-11-188-298-3855	Sequence 3855, App
414	28	53.8	347	11	US-11-087-099-10881	Sequence 10881, A	487	28	53.8	585	11	US-11-188-298-4009	Sequence 4009, App
415	28	53.8	349	11	US-11-096-568a-23821	Sequence 23821, A	488	28	53.8	588	11	US-11-188-298-6479	Sequence 6479, App
416	28	53.8	351	11	US-11-188-298-8113	Sequence 8113, App	489	28	53.8	603	9	US-10-763-712a-95	Sequence 95, App1
417	28	53.8	352	11	US-11-087-099-5907	Sequence 5907, App	490	28	53.8	603	9	US-10-514-581-13	Sequence 8, App1
418	28	53.8	355	11	US-11-087-099-11502	Sequence 11502, A	491	28	53.8	603	9	US-10-514-581-13	Sequence 13, App1
419	28	53.8	356	11	US-11-098-686-10674	Sequence 10674, A	492	28	53.8	635	9	US-10-821-234-1573	Sequence 1573, App
420	28	53.8	356	11	US-11-072-512-3703	Sequence 3703, App	493	28	53.8	637	11	US-11-079-463-5939	Sequence 5939, App
421	28	53.8	360	11	US-11-087-099-9156	Sequence 9156, App	494	28	53.8	686	11	US-11-096-568a-14747	Sequence 14747, A
422	28	53.8	360	11	US-11-188-298-18704	Sequence 18704, A	495	28	53.8	722	9	US-10-469-469-327	Sequence 327, App
423	28	53.8	361	11	US-11-188-298-17456	Sequence 17456, A	496	28	53.8	724	9	US-10-821-234-1506	Sequence 1506, App
424	28	53.8	363	11	US-11-087-099-9277	Sequence 9277, App	497	28	53.8	724	9	US-10-469-469-321	Sequence 321, App
425	28	53.8	363	11	US-11-087-099-10010	Sequence 10010, A	498	28	53.8	724	9	US-10-469-469-322	Sequence 322, App
426	28	53.8	364	9	US-10-703-7998-208	Sequence 208, App	499	28	53.8	724	9	US-10-469-469-326	Sequence 326, App
427	28	53.8	365	11	US-11-096-568a-32809	Sequence 32809, A	500	28	53.8	725	9	US-10-469-469-324	Sequence 324, App
428	28	53.8	367	9	US-10-921-198-62	Sequence 62, App1	501	28	53.8	732	9	US-10-469-469-318	Sequence 318, App
429	28	53.8	367	9	US-10-942-042-62	Sequence 62, App1	502	28	53.8	732	11	US-11-187-230-21	Sequence 21, App1
430	28	53.8	367	9	US-10-519-621-1	Sequence 62, App1	503	28	53.8	733	9	US-10-469-469-323	Sequence 323, App
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432	28	53.8	374	11	US-11-096-568a-33427	Sequence 33427, A	505	28	53.8	758	11	US-11-245-400-30	Sequence 30, App1
433	28	53.8	384	11	US-11-096-568a-33509	Sequence 33509, A	506	28	53.8	759	11	US-11-072-175-167	Sequence 167, App
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435	28	53.8	388	11	US-11-096-568a-33508	Sequence 33508, A	508	28	53.8	772	9	US-10-330-773-373	Sequence 373, App
436	28	53.8	388	11	US-11-188-298-1172	Sequence 1172, App	509	28	53.8	778	11	US-11-188-298-10130	Sequence 10130, A
437	28	53.8	396	11	US-11-096-568a-33507	Sequence 33507, A	510	28	53.8	796	11	US-11-098-686-10952	Sequence 10952, A
438	28	53.8	401	9	US-10-510-386-68	Sequence 68, App1	511	28	53.8	797	9	US-10-469-469-48	Sequence 48, App1
439	28	53.8	405	11	US-11-188-298-5884	Sequence 5884, App	512	28	53.8	797	9	US-10-469-469-208	Sequence 208, App
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441	28	53.8	414	11	US-11-087-099-8301	Sequence 8301, App	514	28	53.8	858	9	US-10-469-469-52	Sequence 52, App1
442	28	53.8	416	9	US-10-469-469-46	Sequence 46, App1	515	28	53.8	858	11	US-11-096-568a-14745	Sequence 14745, A
443	28	53.8	416	9	US-10-469-469-206	Sequence 206, App	516	28	53.8	920	9	US-10-330-773-376	Sequence 376, App
444	28	53.8	419	9	US-10-506-454-989	Sequence 989, App	517	28	53.8	931	8	US-10-505-928-91	Sequence 91, App1
445	28	53.8	428	11	US-11-072-512-3675	Sequence 3675, App	518	28	53.8	1027	9	US-10-330-773-265	Sequence 265, App1
446	28	53.8	432	11	US-11-096-568a-21815	Sequence 21815, A	519	28	53.8	1036	9	US-10-204-673-45	Sequence 45, App1
447	28	53.8	433	11	US-11-087-099-6195	Sequence 6195, App	520	28	53.8	1051	9	US-10-330-773-268	Sequence 268, App
448	28	53.8	433	11	US-11-188-298-5647	Sequence 5647, App	521	28	53.8	1058	9	US-10-821-234-1473	Sequence 1473, App
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450	28	53.8	434	11	US-11-188-298-18319	Sequence 18319, A	523	28	53.8	1058	10	US-11-251-673-3	Sequence 3, App1
451	28	53.8	437	11	US-11-087-099-2535	Sequence 2535, App	524	28	53.8	1058	11	US-11-069-642-105	Sequence 105, App
452	28	53.8	442	11	US-11-096-568a-23131	Sequence 23131, A	525	28	53.8	1071	9	US-10-922-238B-59	Sequence 59, App1
453	28	53.8	449	11	US-11-096-568a-21814	Sequence 21814, A	526	28	53.8	1134	9	US-11-264-096-1570	Sequence 1570, App
454	28	53.8	450	11	US-11-079-463-10401	Sequence 10401, A	527	28	53.8	1185	9	US-10-877-346-7	Sequence 7, App1
455	28	53.8	452	11	US-11-087-099-10811	Sequence 10811, A	528	28	53.8	1419	11	US-11-114-962-3	Sequence 3, App1
456	28	53.8	454	11	US-11-087-099-12184	Sequence 12184, A	529	28	53.8	1451	11	US-11-046-346-1	Sequence 1, App1
457	28	53.8	464	11	US-11-096-568a-30210	Sequence 30210, A	530	28	53.8	1822	8	US-10-505-928-700	Sequence 700, App
458	28	53.8	467	9	US-10-514-581-7	Sequence 7, App1	531	28	53.8	1822	11	US-11-169-041-193	Sequence 193, App
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533	28	53.8	2117	11	US-11-188-298-8910	Sequence 8910, Ap	606	27	51.9	218	11	US-11-096-568A-1358	Sequence 1358, Ap
534	28	53.8	2132	11	US-11-087-099-2434	Sequence 2434, Ap	607	27	51.9	221	9	US-10-821-234-1139	Sequence 1139, Ap
535	28	53.8	2132	11	US-11-188-298-2344	Sequence 2344, Ap	608	27	51.9	226	11	US-11-132-283-5	Sequence 5, Appl1
536	28	53.8	2471	11	US-11-050-346-68	Sequence 68, Appl	609	27	51.9	226	11	US-11-096-568A-19476	Sequence 19476, A
537	28	53.8	4060	9	US-10-922-2328-55	Sequence 55, Appl	610	27	51.9	228	11	US-11-096-568A-19475	Sequence 19475, A
538	28	53.8	4386	11	US-11-004-399-714	Sequence 714, Ap	611	27	51.9	231	11	US-11-132-283-61	Sequence 61, Appl
539	28	53.8	6738	9	US-10-922-2328-56	Sequence 56, Appl	612	27	51.9	233	11	US-11-096-568A-10916	Sequence 10916, A
540	28	53.8	8746	11	US-11-098-686-10232	Sequence 10232, A	613	27	51.9	235	11	US-11-087-099-5329	Sequence 5329, Ap
541	27.5	52.9	258	11	US-11-188-298-4937	Sequence 4937, Ap	614	27	51.9	235	11	US-11-096-568A-1357	Sequence 1357, Ap
542	27	51.9	9	9	US-10-530-061-629	Sequence 629, App	615	27	51.9	235	11	US-11-096-568A-10915	Sequence 10915, A
543	27	51.9	9	9	US-10-530-061-630	Sequence 630, App	616	27	51.9	239	11	US-11-072-512-2162	Sequence 2162, Ap
544	27	51.9	10	11	US-11-004-399-778	Sequence 778, App	617	27	51.9	240	11	US-11-188-298-13805	Sequence 13805, A
545	27	51.9	34	11	US-11-004-399-1830	Sequence 1830, Ap	618	27	51.9	241	11	US-11-096-568A-5617	Sequence 5617, Ap
546	27	51.9	37	9	US-10-467-657-1382	Sequence 1382, Ap	619	27	51.9	246	11	US-11-096-568A-10949	Sequence 10949, A
547	27	51.9	54	9	US-10-467-657-3784	Sequence 3784, Ap	620	27	51.9	246	11	US-11-096-568A-19474	Sequence 19474, A
548	27	51.9	68	9	US-10-467-657-9122	Sequence 9122, Ap	621	27	51.9	250	11	US-11-087-099-9977	Sequence 9977, Ap
549	27	51.9	77	9	US-10-925-366A-305	Sequence 305, App	622	27	51.9	251	11	US-11-054-515-210	Sequence 210, App
550	27	51.9	77	11	US-11-172-740-2391	Sequence 2391, Ap	623	27	51.9	251	11	US-11-072-512-3719	Sequence 3719, Ap
551	27	51.9	77	11	US-11-098-758-305	Sequence 305, App	624	27	51.9	251	11	US-11-072-512-3719	Sequence 3719, Ap
552	27	51.9	81	11	US-11-096-568A-3707	Sequence 3707, Ap	625	27	51.9	251	11	US-11-266-444-210	Sequence 22079, A
553	27	51.9	88	11	US-11-045-004-2443	Sequence 2443, Ap	626	27	51.9	261	11	US-11-079-463-9081	Sequence 9081, Ap
554	27	51.9	90	11	US-11-264-096-76	Sequence 76, Appl	627	27	51.9	262	11	US-11-172-740-2394	Sequence 2394, Ap
555	27	51.9	92	11	US-11-264-096-1410	Sequence 1410, Ap	628	27	51.9	277	11	US-11-072-512-2132	Sequence 2132, Ap
556	27	51.9	93	9	US-10-506-454-417	Sequence 417, App	629	27	51.9	288	11	US-11-079-463-6303	Sequence 6303, Ap
557	27	51.9	100	11	US-11-096-568A-6932	Sequence 6932, Ap	630	27	51.9	297	9	US-10-967-527A-17	Sequence 17, Appl
558	27	51.9	107	11	US-11-217-819-114	Sequence 114, App	631	27	51.9	307	11	US-11-096-568A-6261	Sequence 6261, Ap
559	27	51.9	108	9	US-10-925-366A-200	Sequence 200, App	632	27	51.9	314	11	US-11-096-568A-10948	Sequence 10948, A
560	27	51.9	108	9	US-10-925-366A-207	Sequence 207, App	633	27	51.9	315	11	US-11-087-099-1391	Sequence 1391, Ap
561	27	51.9	108	11	US-11-098-758-200	Sequence 200, App	634	27	51.9	315	11	US-11-087-099-4663	Sequence 4663, Ap
562	27	51.9	108	11	US-11-098-758-207	Sequence 207, App	635	27	51.9	315	11	US-11-087-099-5729	Sequence 5729, Ap
563	27	51.9	112	11	US-11-172-740-2390	Sequence 2390, Ap	636	27	51.9	315	11	US-11-087-099-7813	Sequence 7813, Ap
564	27	51.9	123	9	US-10-508-307-3	Sequence 3, Appl1	637	27	51.9	315	11	US-11-087-099-7956	Sequence 7956, Ap
565	27	51.9	127	11	US-11-096-568A-3706	Sequence 3706, Ap	638	27	51.9	315	11	US-11-087-099-8869	Sequence 8869, Ap
566	27	51.9	131	11	US-11-087-099-6699	Sequence 6699, App	639	27	51.9	315	11	US-11-087-099-12243	Sequence 12243, A
567	27	51.9	131	11	US-11-172-740-735	Sequence 735, App	640	27	51.9	315	11	US-11-096-568A-2781	Sequence 2781, Ap
568	27	51.9	132	11	US-11-087-099-7442	Sequence 7442, Ap	641	27	51.9	317	11	US-11-096-568A-6260	Sequence 6260, Ap
569	27	51.9	142	9	US-10-467-657-1558	Sequence 1558, Ap	642	27	51.9	318	11	US-11-096-568A-23270	Sequence 23270, A
570	27	51.9	153	11	US-11-079-463-6445	Sequence 6445, Ap	643	27	51.9	319	11	US-11-087-099-8640	Sequence 8640, Ap
571	27	51.9	157	11	US-11-072-512-2634	Sequence 2634, Ap	644	27	51.9	320	11	US-11-096-568A-1118	Sequence 1218, Ap
572	27	51.9	158	11	US-11-087-099-7882	Sequence 7882, Ap	645	27	51.9	322	11	US-11-098-686-10665	Sequence 10665, A
573	27	51.9	159	11	US-11-132-285-7	Sequence 7, Appl1	646	27	51.9	324	11	US-11-096-568A-23269	Sequence 23269, A
574	27	51.9	163	11	US-11-079-463-6072	Sequence 6072, Ap	647	27	51.9	325	11	US-11-087-099-23268	Sequence 23268, A
575	27	51.9	163	11	US-11-174-121-4	Sequence 4, Appl1	648	27	51.9	330	11	US-11-087-099-2760	Sequence 2760, Ap
576	27	51.9	182	11	US-11-079-463-7021	Sequence 5, Appl1	649	27	51.9	340	11	US-11-096-568A-22078	Sequence 22078, A
577	27	51.9	182	11	US-11-188-298-7966	Sequence 7966, Ap	650	27	51.9	341	11	US-11-087-099-10534	Sequence 10534, A
578	27	51.9	182	11	US-11-188-298-7966	Sequence 7966, Ap	651	27	51.9	342	11	US-11-096-568A-2780	Sequence 2780, Ap
579	27	51.9	195	11	US-11-096-568A-9232	Sequence 9232, Ap	652	27	51.9	346	11	US-11-098-686-10637	Sequence 10637, A
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581	27	51.9	197	11	US-11-087-099-618	Sequence 618, App	654	27	51.9	349	11	US-11-087-099-3333	Sequence 3333, Ap
582	27	51.9	197	11	US-11-087-099-830	Sequence 830, App	655	27	51.9	350	11	US-11-087-099-2437	Sequence 2437, Ap
583	27	51.9	197	11	US-11-087-099-1498	Sequence 1498, Ap	656	27	51.9	350	11	US-11-087-099-1952	Sequence 1952, A
584	27	51.9	197	11	US-11-087-099-2027	Sequence 2027, Ap	657	27	51.9	350	11	US-11-087-099-12657	Sequence 12657, Ap
585	27	51.9	197	11	US-11-087-099-2599	Sequence 2599, Ap	658	27	51.9	355	9	US-10-506-454-140	Sequence 340, App
586	27	51.9	197	11	US-11-087-099-2970	Sequence 2970, Ap	659	27	51.9	355	11	US-11-188-298-17522	Sequence 17522, A
587	27	51.9	197	11	US-11-087-099-3199	Sequence 3199, Ap	660	27	51.9	358	11	US-11-087-099-10697	Sequence 10697, A
588	27	51.9	197	11	US-11-087-099-3824	Sequence 3824, Ap	661	27	51.9	359	11	US-11-087-099-4199	Sequence 4199, Ap
589	27	51.9	197	11	US-11-087-099-4056	Sequence 4056, Ap	662	27	51.9	359	11	US-11-087-099-6932	Sequence 6932, Ap
590	27	51.9	197	11	US-11-087-099-4723	Sequence 4723, Ap	663	27	51.9	359	11	US-11-172-740-2393	Sequence 2393, Ap
591	27	51.9	197	11	US-11-087-099-4781	Sequence 4781, Ap	664	27	51.9	364	11	US-11-188-298-20895	Sequence 20895, A
592	27	51.9	197	11	US-11-087-099-6230	Sequence 6230, Ap	665	27	51.9	365	11	US-11-072-512-3183	Sequence 3183, Ap
593	27	51.9	197	11	US-11-087-099-6469	Sequence 6469, Ap	666	27	51.9	364	11	US-11-188-298-19989	Sequence 19989, A
594	27	51.9	197	11	US-11-087-099-7877	Sequence 7877, Ap	667	27	51.9	364	11	US-11-188-298-13415	Sequence 13415, A
595	27	51.9	197	11	US-11-087-099-9396	Sequence 9396, Ap	668	27	51.9	369	11	US-11-087-099-2966	Sequence 2966, Ap
596	27	51.9	197	11	US-11-087-099-10338	Sequence 10338, A	669	27	51.9	369	11	US-11-087-099-2978	Sequence 2978, Ap
597	27	51.9	197	11	US-11-087-099-11110	Sequence 11110, A	670	27	51.9	371	11	US-11-186-284-16	Sequence 16, Appl
598	27	51.9	197	11	US-11-087-099-11403	Sequence 11403, A	671	27	51.9	375	9	US-10-793-626-2918	Sequence 2918, Ap
599	27	51.9	197	11	US-11-087-099-11974	Sequence 11974, A	672	27	51.9	375	11	US-11-188-298-2280	Sequence 2280, Ap
600	27	51.9	202	9	US-10-467-657-8154	Sequence 8154, Ap	673	27	51.9	376	11	US-11-087-099-2815	Sequence 2815, Ap
601	27	51.9	205	11	US-11-096-568A-10917	Sequence 10917, A	674	27	51.9	378	11	US-11-188-298-16585	Sequence 16585, A
602	27	51.9	208	11	US-11-096-568A-9230	Sequence 9230, Ap	675	27	51.9	378	11	US-11-188-298-17787	Sequence 17787, A
603	27	51.9	212	11	US-11-172-740-2389	Sequence 2389, Ap	676	27	51.9	379	11	US-11-087-099-1863	Sequence 1863, Ap
604	27	51.9	213	11	US-11-096-568A-5618	Sequence 5618, Ap	677	27	51.9	379	11	US-11-087-099-1863	Sequence 1863, Ap
605	27	51.9	215	11	US-11-264-096-2096	Sequence 2096, Ap	678	27	51.9	379	11	US-11-096-568A-11269	Sequence 11269, A

679	27	51.9	380	11	US-11-087-099-4525	Sequence 4525, Ap	752	27	51.9	537	9	US-10-501-035-318	Sequence 318, App
680	27	51.9	380	11	US-11-087-099-4810	Sequence 4810, Ap	753	27	51.9	541	11	US-11-166-412-55	Sequence 55, App1
681	27	51.9	380	11	US-11-087-099-5011	Sequence 5011, Ap	754	27	51.9	546	11	US-11-040-218-15	Sequence 15, App1
682	27	51.9	380	11	US-11-087-099-6231	Sequence 6231, Ap	755	27	51.9	553	11	US-11-188-298-6247	Sequence 6247, Ap
683	27	51.9	380	11	US-11-087-099-8371	Sequence 8371, Ap	756	27	51.9	558	9	US-10-467-657-4258	Sequence 4258, Ap
684	27	51.9	380	11	US-11-087-099-8656	Sequence 8656, Ap	757	27	51.9	564	11	US-11-087-099-3568	Sequence 3568, Ap
685	27	51.9	380	11	US-11-096-568A-1268	Sequence 1268, A	758	27	51.9	568	11	US-11-188-298-8765	Sequence 8765, Ap
686	27	51.9	380	11	US-11-188-298-3577	Sequence 3577, Ap	759	27	51.9	569	11	US-11-188-298-494	Sequence 494, App
687	27	51.9	380	11	US-11-188-298-4130	Sequence 4190, Ap	760	27	51.9	572	9	US-10-821-234-1290	Sequence 1290, Ap
688	27	51.9	380	11	US-11-188-298-4411	Sequence 4411, Ap	761	27	51.9	575	11	US-11-072-175-201	Sequence 2015, App
689	27	51.9	380	11	US-11-172-740-2388	Sequence 2388, Ap	762	27	51.9	578	11	US-11-188-298-20155	Sequence 17, App1
690	27	51.9	385	11	US-11-172-740-2387	Sequence 2387, Ap	763	27	51.9	582	8	US-10-505-928-813	Sequence 813, App
691	27	51.9	390	9	US-10-506-454-1085	Sequence 1085, Ap	764	27	51.9	582	11	US-11-090-439-58	Sequence 58, App1
692	27	51.9	392	11	US-11-096-568A-1217	Sequence 1217, Ap	765	27	51.9	582	11	US-11-169-041-130	Sequence 130, App
693	27	51.9	394	11	US-11-087-099-11627	Sequence 11627, Ap	766	27	51.9	582	11	US-11-1200-822-2	Sequence 2, App1
694	27	51.9	403	9	US-10-485-517-165	Sequence 165, App	767	27	51.9	582	11	US-11-087-099-7593	Sequence 7593, Ap
695	27	51.9	404	11	US-11-096-568A-22077	Sequence 22077, A	768	27	51.9	585	11	US-11-188-298-20009	Sequence 20009, A
696	27	51.9	407	11	US-11-096-568A-2779	Sequence 2779, Ap	769	27	51.9	587	11	US-11-188-298-472	Sequence 472, App
697	27	51.9	407	11	US-11-096-568A-2782	Sequence 2782, Ap	770	27	51.9	590	11	US-11-040-218-11	Sequence 11, App1
698	27	51.9	409	11	US-11-096-568A-10947	Sequence 10947, A	771	27	51.9	598	11	US-11-198-640A-2	Sequence 2, App1
699	27	51.9	410	8	US-10-505-928-692	Sequence 692, App	772	27	51.9	598	11	US-11-188-298-3661	Sequence 3661, Ap
700	27	51.9	410	11	US-11-079-463-5336	Sequence 5336, Ap	773	27	51.9	611	11	US-11-188-298-8258	Sequence 20983, A
701	27	51.9	413	11	US-11-096-568A-11267	Sequence 11267, A	774	27	51.9	616	11	US-11-188-298-8258	Sequence 8258, Ap
702	27	51.9	417	9	US-10-506-454-424	Sequence 424, App	775	27	51.9	632	11	US-11-096-568A-33996	Sequence 33996, Ap
703	27	51.9	423	9	US-10-506-454-1080	Sequence 1080, Ap	776	27	51.9	645	9	US-10-821-234-1409	Sequence 1409, Ap
704	27	51.9	432	11	US-11-172-740-2396	Sequence 2396, Ap	777	27	51.9	662	11	US-11-072-512-3349	Sequence 3349, Ap
705	27	51.9	433	11	US-11-245-400-19	Sequence 19, App1	778	27	51.9	666	9	US-10-745-586-188	Sequence 188, App
706	27	51.9	438	11	US-11-096-568A-6259	Sequence 6259, Ap	779	27	51.9	668	11	US-11-106-674-11	Sequence 1, App1
707	27	51.9	439	11	US-11-096-568A-1216	Sequence 1216, Ap	780	27	51.9	699	11	US-11-188-298-21163	Sequence 21163, A
708	27	51.9	441	10	US-11-242-111-29	Sequence 29, App1	781	27	51.9	696	11	US-11-188-298-13976	Sequence 13976, A
709	27	51.9	444	9	US-10-467-657-2414	Sequence 2414, Ap	782	27	51.9	702	11	US-11-098-686-10194	Sequence 10194, A
710	27	51.9	445	9	US-10-453-372-2	Sequence 2, App1	783	27	51.9	808	11	US-11-072-512-2819	Sequence 2819, Ap
711	27	51.9	449	11	US-11-087-099-7707	Sequence 7707, Ap	784	27	51.9	808	9	US-10-330-773-195	Sequence 195, App
712	27	51.9	450	11	US-11-186-284-14	Sequence 14, App1	785	27	51.9	906	9	US-10-329-258-12	Sequence 12, App1
713	27	51.9	451	11	US-11-188-298-13702	Sequence 13702, A	786	27	51.9	968	9	US-10-821-234-1188	Sequence 1188, App
714	27	51.9	453	11	US-11-087-099-6300	Sequence 6300, Ap	787	27	51.9	968	11	US-11-096-568A-11193	Sequence 31193, A
715	27	51.9	453	11	US-11-188-298-16793	Sequence 16793, A	788	27	51.9	994	11	US-11-096-568A-11192	Sequence 31191, A
716	27	51.9	454	11	US-11-098-686-10919	Sequence 10919, A	789	27	51.9	994	11	US-11-096-568A-11191	Sequence 31191, A
717	27	51.9	457	11	US-11-087-099-1111	Sequence 1111, Ap	790	27	51.9	1048	9	US-10-501-035-229	Sequence 229, App
718	27	51.9	457	11	US-11-188-298-12060	Sequence 12060, A	791	27	51.9	1050	8	US-10-505-928-347	Sequence 347, App
719	27	51.9	462	11	US-11-087-099-7509	Sequence 7509, Ap	792	27	51.9	1050	9	US-10-523-477-12	Sequence 12, App1
720	27	51.9	462	11	US-11-188-298-17974	Sequence 17974, A	793	27	51.9	1059	9	US-10-770-726-47	Sequence 47, App1
721	27	51.9	464	11	US-11-188-298-8888	Sequence 8888, Ap	794	27	51.9	1059	11	US-11-098-686-10296	Sequence 10296, A
722	27	51.9	466	11	US-11-087-099-10242	Sequence 10242, A	795	27	51.9	1059	9	US-10-469-469-91	Sequence 91, App1
723	27	51.9	469	11	US-11-188-298-2971	Sequence 2971, Ap	796	27	51.9	1104	9	US-10-330-773-794	Sequence 794, App
724	27	51.9	469	11	US-11-188-298-9435	Sequence 9435, Ap	797	27	51.9	1112	11	US-11-096-568A-33821	Sequence 33820, A
725	27	51.9	471	11	US-11-072-512-2636	Sequence 2636, Ap	798	27	51.9	1113	11	US-11-096-568A-33820	Sequence 33819, A
726	27	51.9	475	8	US-10-505-928-142	Sequence 142, App	799	27	51.9	1115	11	US-11-096-568A-33819	Sequence 190, App
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728	27	51.9	484	9	US-10-763-712A-12	Sequence 12, App1	801	27	51.9	1202	9	US-10-330-773-193	Sequence 28513, Ap
729	27	51.9	485	11	US-11-087-099-1672	Sequence 1672, Ap	802	27	51.9	1233	11	US-11-096-568A-18513	Sequence 28512, A
730	27	51.9	489	11	US-11-096-568A-33382	Sequence 32382, A	803	27	51.9	1401	11	US-11-096-568A-18512	Sequence 1, App1
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732	27	51.9	492	11	US-11-072-512-1974	Sequence 1974, Ap	805	27	51.9	1659	9	US-11-072-175-205	Sequence 10177, A
733	27	51.9	492	11	US-11-072-512-3223	Sequence 3223, Ap	806	27	51.9	1659	11	US-11-096-568A-10177	Sequence 10175, A
734	27	51.9	495	11	US-11-166-412-54	Sequence 54, App1	807	26.5	51.0	196	11	US-11-096-568A-10175	Sequence 10175, A
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736	27	51.9	499	11	US-11-087-099-10769	Sequence 10769, A	809	26.5	51.0	424	11	US-11-087-099-3416	Sequence 3416, Ap
737	27	51.9	499	11	US-11-188-298-9952	Sequence 9952, Ap	810	26	50.0	9	US-10-530-061-807	Sequence 807, App	
738	27	51.9	500	11	US-11-166-412-53	Sequence 53, App1	811	26	50.0	10	US-10-505-955-67	Sequence 807, App	
739	27	51.9	500	11	US-11-040-218-13	Sequence 13, App1	812	26	50.0	13	US-10-895-064-382	Sequence 382, App	
740	27	51.9	503	11	US-11-087-099-9776	Sequence 9776, Ap	813	26	50.0	13	US-11-129-741-382	Sequence 3312, Ap	
741	27	51.9	503	11	US-11-188-298-20070	Sequence 20070, A	814	26	50.0	13	US-11-129-741-382	Sequence 269, App	
742	27	51.9	505	9	US-10-519-447-4	Sequence 4, App1	815	26	50.0	13	US-11-152-974A-182	Sequence 182, App	
743	27	51.9	505	11	US-11-242-111-28	Sequence 28, App1	816	26	50.0	17	US-11-152-974A-182	Sequence 182, App	
744	27	51.9	505	11	US-11-188-298-8716	Sequence 8716, Ap	817	26	50.0	17	US-11-152-974A-182	Sequence 182, App	
745	27	51.9	505	11	US-11-188-298-21337	Sequence 21337, A	818	26	50.0	17	US-11-153-143A-183	Sequence 183, App	
746	27	51.9	506	11	US-11-188-298-9019	Sequence 9019, Ap	819	26	50.0	17	US-11-153-143A-183	Sequence 183, App	
747	27	51.9	506	11	US-11-188-298-9318	Sequence 9318, Ap	820	26	50.0	19	US-11-153-143A-183	Sequence 183, App	
748	27	51.9	523	11	US-11-072-512-2046	Sequence 2046, Ap	821	26	50.0	19	US-11-153-143A-550	Sequence 550, App	
749	27	51.9	532	11	US-11-040-218-19	Sequence 19, App1	822	26	50.0	20	US-11-004-399-1818	Sequence 1818, Ap	
750	27	51.9	532	11	US-11-079-463-5528	Sequence 5528, Ap	823	26	50.0	22	9	US-10-467-657-6908	Sequence 6908, Ap
751	27	51.9	537	9	US-10-912-971-14	Sequence 14, App1	824	26	50.0	22	9	US-10-467-657-6908	Sequence 6908, Ap

825	26	50.0	23	11	US-11-236-657-225	Sequence 225, App	898	26	50.0	158	11	US-11-096-568A-28648	Sequence 28648, A
826	26	50.0	29	11	US-11-207-078-162	Sequence 162, App	899	26	50.0	160	9	US-10-467-657-6486	Sequence 6486, App
827	26	50.0	32	11	US-11-004-399-1201	Sequence 1201, App	900	26	50.0	160	11	US-11-010-874-12	Sequence 12, App
828	26	50.0	35	9	US-10-957-351-10	Sequence 10, App	901	26	50.0	160	11	US-11-124-368A-332	Sequence 332, App
829	26	50.0	36	9	US-10-485-517-138	Sequence 138, App	902	26	50.0	161	9	US-10-511-538-251	Sequence 251, App
830	26	50.0	41	11	US-11-264-096-1023	Sequence 1023, App	903	26	50.0	161	11	US-11-154-257-3	Sequence 3, App
831	26	50.0	41	11	US-11-264-096-1024	Sequence 1024, App	904	26	50.0	163	11	US-11-079-463-8934	Sequence 8934, App
832	26	50.0	45	11	US-11-226-657-71	Sequence 71, App	905	26	50.0	167	11	US-11-096-568A-14802	Sequence 14802, A
833	26	50.0	46	9	US-10-895-064-1666	Sequence 1666, App	906	26	50.0	167	11	US-11-264-096-2217	Sequence 2217, App
834	26	50.0	46	11	US-11-129-741-1666	Sequence 1666, App	907	26	50.0	168	11	US-11-010-874-7	Sequence 7, App
835	26	50.0	46	11	US-11-096-568A-2613	Sequence 2613, App	908	26	50.0	168	11	US-11-079-463-5900	Sequence 5900, App
836	26	50.0	46	11	US-11-004-399-360	Sequence 360, App	909	26	50.0	168	11	US-11-188-298-114025	Sequence 14025, A
837	26	50.0	46	11	US-11-004-399-691	Sequence 691, App	910	26	50.0	171	11	US-11-188-298-11096	Sequence 11096, A
838	26	50.0	46	11	US-11-004-399-3721	Sequence 3721, App	911	26	50.0	173	11	US-11-096-568A-20289	Sequence 20289, A
839	26	50.0	51	9	US-10-986-501-131	Sequence 131, App	912	26	50.0	179	11	US-11-188-298-21944	Sequence 21944, A
840	26	50.0	56	9	US-10-475-075-885	Sequence 885, App	913	26	50.0	180	11	US-11-079-463-8232	Sequence 8262, App
841	26	50.0	58	11	US-11-004-399-2282	Sequence 2282, App	914	26	50.0	180	11	US-11-188-298-11609	Sequence 1609, App
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845	26	50.0	75	9	US-10-467-657-4938	Sequence 4938, App	918	26	50.0	186	11	US-11-096-568A-28647	Sequence 28647, A
846	26	50.0	80	9	US-10-131-826A-84	Sequence 84, App	919	26	50.0	189	11	US-11-188-298-13424	Sequence 13424, A
847	26	50.0	80	9	US-10-973-115B-84	Sequence 84, App	920	26	50.0	192	11	US-11-067-527A-28	Sequence 28, App
848	26	50.0	80	9	US-10-137-873A-84	Sequence 84, App	921	26	50.0	193	9	US-10-967-703-142	Sequence 142, App
849	26	50.0	80	9	US-10-152-370-84	Sequence 84, App	922	26	50.0	193	11	US-11-102-240-142	Sequence 142, App
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851	26	50.0	83	11	US-11-179-844-3	Sequence 3, App	924	26	50.0	194	11	US-11-172-740-2028	Sequence 2028, App
852	26	50.0	86	9	US-10-469-469-237	Sequence 237, App	925	26	50.0	197	11	US-11-098-686-11017	Sequence 11017, A
853	26	50.0	87	11	US-11-226-657-221	Sequence 221, App	926	26	50.0	198	9	US-10-469-469-339	Sequence 339, App
854	26	50.0	93	11	US-11-188-298-17895	Sequence 17895, A	927	26	50.0	198	9	US-10-469-469-241	Sequence 241, App
855	26	50.0	94	9	US-10-821-234-1066	Sequence 1066, App	928	26	50.0	201	11	US-11-188-298-8986	Sequence 8986, App
856	26	50.0	95	11	US-11-004-590-74	Sequence 74, App	929	26	50.0	202	11	US-11-079-463-8895	Sequence 8895, App
857	26	50.0	95	11	US-11-096-568A-3962	Sequence 3962, App	930	26	50.0	205	11	US-11-172-740-2025	Sequence 2025, App
858	26	50.0	96	11	US-11-084-554-132	Sequence 132, App	931	26	50.0	207	11	US-11-096-568A-4691	Sequence 4691, App
859	26	50.0	96	11	US-11-136-250-132	Sequence 132, App	932	26	50.0	209	11	US-11-045-004-149	Sequence 149, App
860	26	50.0	97	9	US-10-530-253-29	Sequence 29, App	933	26	50.0	210	11	US-11-122-296-21	Sequence 21, App
861	26	50.0	98	8	US-10-511-814-8	Sequence 8, App	934	26	50.0	210	11	US-11-188-298-8720	Sequence 8720, App
862	26	50.0	98	8	US-10-511-814-11	Sequence 11, App	935	26	50.0	211	11	US-11-096-568A-20287	Sequence 20287, A
863	26	50.0	98	9	US-10-530-253-14	Sequence 14, App	936	26	50.0	212	11	US-11-096-568A-20287	Sequence 20287, A
864	26	50.0	98	11	US-11-179-478A-4	Sequence 4, App	937	26	50.0	212	11	US-11-079-463-5397	Sequence 5397, App
865	26	50.0	99	9	US-10-530-253-30	Sequence 30, App	938	26	50.0	213	11	US-11-079-463-9239	Sequence 9239, App
866	26	50.0	104	11	US-11-179-844-1	Sequence 1, App	939	26	50.0	214	11	US-10-821-234-1544	Sequence 1544, App
867	26	50.0	104	11	US-11-284-905-14	Sequence 14, App	940	26	50.0	215	9	US-10-821-234-1544	Sequence 1544, App
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869	26	50.0	110	11	US-11-206-587-30	Sequence 30, App	942	26	50.0	218	9	US-10-374-954-5	Sequence 5, App
870	26	50.0	111	9	US-10-523-362-26	Sequence 26, App	943	26	50.0	218	10	US-11-263-326-129	Sequence 129, App
871	26	50.0	112	11	US-11-172-740-3027	Sequence 3027, App	944	26	50.0	218	10	US-11-263-326-130	Sequence 130, App
872	26	50.0	115	11	US-11-096-568A-1968	Sequence 1968, App	945	26	50.0	218	10	US-11-263-326-174	Sequence 174, App
873	26	50.0	116	11	US-11-051-720-1522	Sequence 1522, App	946	26	50.0	218	11	US-11-158-505-4	Sequence 4, App
874	26	50.0	117	11	US-11-079-463-8766	Sequence 8766, App	947	26	50.0	218	11	US-11-158-505-12	Sequence 12, App
875	26	50.0	117	11	US-11-188-298-22298	Sequence 22298, A	948	26	50.0	218	11	US-11-158-505-10	Sequence 10, App
876	26	50.0	118	11	US-11-090-439-3	Sequence 3, App	949	26	50.0	218	11	US-11-158-505-28	Sequence 28, App
877	26	50.0	118	11	US-11-033-039-531	Sequence 531, App	950	26	50.0	220	11	US-11-188-298-14601	Sequence 14601, A
878	26	50.0	118	11	US-11-135-288-2	Sequence 2, App	951	26	50.0	220	11	US-11-096-568A-2809	Sequence 2809, App
879	26	50.0	120	9	US-10-523-362-16	Sequence 16, App	952	26	50.0	222	11	US-11-264-096-1141	Sequence 1141, App
880	26	50.0	124	9	US-10-218-784-28	Sequence 28, App	953	26	50.0	225	11	US-11-158-505-3	Sequence 3, App
881	26	50.0	124	9	US-10-219-061-28	Sequence 28, App	954	26	50.0	227	11	US-11-158-505-11	Sequence 11, App
882	26	50.0	124	9	US-10-219-062-28	Sequence 28, App	955	26	50.0	229	11	US-11-079-463-7328	Sequence 7328, A
883	26	50.0	124	9	US-10-219-064-28	Sequence 28, App	956	26	50.0	230	11	US-11-096-568A-21243	Sequence 21243, A
884	26	50.0	124	9	US-10-333-134-28	Sequence 28, App	957	26	50.0	234	11	US-11-188-298-4916	Sequence 4916, App
885	26	50.0	127	9	US-10-986-501-330	Sequence 330, App	958	26	50.0	236	11	US-11-188-298-16088	Sequence 16088, A
886	26	50.0	133	9	US-10-523-362-22	Sequence 22, App	959	26	50.0	238	11	US-11-158-505-1	Sequence 1, App
887	26	50.0	133	11	US-11-096-568-10388	Sequence 10388, A	960	26	50.0	238	11	US-11-158-505-3	Sequence 3, App
888	26	50.0	136	9	US-10-523-362-46	Sequence 46, App	961	26	50.0	238	11	US-11-158-505-17	Sequence 17, App
889	26	50.0	139	8	US-10-505-928-179	Sequence 179, App	962	26	50.0	238	11	US-11-158-505-19	Sequence 19, App
890	26	50.0	142	10	US-11-254-182-22	Sequence 22, App	963	26	50.0	238	11	US-11-158-505-25	Sequence 25, App
891	26	50.0	142	11	US-11-182-908-22	Sequence 22, App	964	26	50.0	238	11	US-11-158-505-27	Sequence 27, App
892	26	50.0	143	11	US-11-079-463-10266	Sequence 10266, A	965	26	50.0	238	11	US-11-158-505-7	Sequence 7, App
893	26	50.0	145	11	US-11-156-084-130	Sequence 130, App	967	26	50.0	238	11	US-11-158-505-74	Sequence 74, App
894	26	50.0	149	9	US-10-213-292-40	Sequence 40, App	968	26	50.0	239	9	US-11-264-096-1383	Sequence 1383, App
895	26	50.0	152	11	US-11-188-298-2828	Sequence 2828, App	969	26	50.0	240	11	US-10-980-388-108	Sequence 108, App
896	26	50.0	152	11	US-11-188-298-3999	Sequence 3999, App	970	26	50.0	240	11	US-11-096-568A-13122	Sequence 13122, A
897	26	50.0	157	11	US-11-172-740-2471	Sequence 2471, App							

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971 26 50.0 241 11 US-11-059-275-2 Sequence 2, Appli
972 26 50.0 241 11 US-11-079-463-9266 Sequence 9266, Ap
973 26 50.0 242 11 US-11-096-568A-13121 Sequence 13121, A
974 26 50.0 247 11 US-11-188-298-11688 Sequence 11688, A
975 26 50.0 248 11 US-11-188-298-11108 Sequence 11108, A
976 26 50.0 251 9 US-10-485-517-167 Sequence 167, App
977 26 50.0 251 11 US-11-079-463-10128 Sequence 10128, A
978 26 50.0 252 9 US-10-506-454-818 Sequence 818, App
979 26 50.0 252 11 US-11-096-568A-21242 Sequence 21242, A
980 26 50.0 253 11 US-11-045-004-745 Sequence 745, App
981 26 50.0 255 11 US-11-188-298-8279 Sequence 8279, Ap
982 26 50.0 255 11 US-11-188-298-21413 Sequence 21413, A
983 26 50.0 258 11 US-11-188-298-2657 Sequence 2657, Ap
984 26 50.0 259 11 US-11-045-004-366 Sequence 366, App
985 26 50.0 263 11 US-11-072-512-3558 Sequence 3558, Ap
986 26 50.0 264 11 US-11-096-568A-5759 Sequence 5759, Ap
987 26 50.0 268 9 US-10-469-469-342 Sequence 242, App
988 26 50.0 268 11 US-11-096-568A-22646 Sequence 22646, A
989 26 50.0 270 11 US-11-096-568A-21566 Sequence 21566, A
990 26 50.0 272 11 US-11-046-456-23 Sequence 23, Appl
991 26 50.0 272 11 US-11-046-644-23 Sequence 23, Appl
992 26 50.0 272 11 US-11-137-253-14 Sequence 14, Appl
993 26 50.0 273 11 US-11-188-298-3841 Sequence 3841, Ap
994 26 50.0 278 11 US-11-096-568A-2808 Sequence 2808, Ap
995 26 50.0 281 11 US-11-188-298-7735 Sequence 7735, Ap
996 26 50.0 282 11 US-11-264-096-277 Sequence 277, App
997 26 50.0 283 11 US-11-096-568A-29222 Sequence 29222, A
998 26 50.0 290 11 US-11-096-568A-20725 Sequence 20725, A
999 26 50.0 291 9 US-10-467-657-6508 Sequence 6508, Ap
1000 26 50.0 293 11 US-11-096-568A-32227 Sequence 32227, A
```

ALIGNMENTS

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RESULT 1
US-10-530-253-13
; Sequence 13, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 13
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-13
Query Match 100.0%; Score 52; DB 9; Length 151;
Best Local Similarity 100.0%; Pred. No. 0.04;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 30 CVYCKQQL 38
```

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RESULT 2
US-11-206-138-3
; Sequence 3, Application US/11206138
; Publication No. US20060039919A1
```

```
; GENERAL INFORMATION:
; APPLICANT: Healthbancs Biotech CO. LTD.
; TITLE OF INVENTION: Fusion protein for inhibiting cervical cancer
; FILE REFERENCE: P7819/0613
; CURRENT APPLICATION NUMBER: US/11/206,138
; CURRENT FILING DATE: 2005-08-18
; NUMBER OF SEQ ID NOS: 14
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 3
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-11-206-138-3
```

```
Query Match 100.0%; Score 52; DB 11; Length 158;
Best Local Similarity 100.0%; Pred. No. 0.041;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
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```
QY 1 CVYCKQQL 9
Db 37 CVYCKQQL 45
```

```
RESULT 3
US-10-530-253-1
; Sequence 1, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 1
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-1
```

```
Query Match 100.0%; Score 52; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQQL 9
Db 30 CVYCKQQL 38
```

```
RESULT 4
US-10-530-253-3
; Sequence 3, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Caesetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
```

```
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 3
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-3
```

```
Query Match          100.0%; Score 52; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQQL 9
        |||||
Db       30 CVYCKQQL 38
```

```
RESULT 5
US-10-530-253-5
; Sequence 5, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-5
```

```
Query Match          100.0%; Score 52; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQQL 9
        |||||
Db       30 CVYCKQQL 38
```

```
RESULT 6
US-10-530-253-7
; Sequence 7, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 248
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```
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-7
```

```
-Query Match          100.0%; Score 52; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQQL 9
        |||||
Db       127 CVYCKQQL 135
```

```
RESULT 7
US-10-530-253-9
; Sequence 9, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 9
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-9
```

```
Query Match          100.0%; Score 52; DB 9; Length 248;
Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQQL 9
        |||||
Db       127 CVYCKQQL 135
```

```
RESULT 8
US-10-530-253-11
; Sequence 11, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casaretti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT FILING DATE: 2005-04-04
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 11
; LENGTH: 248
; TYPE: PRT
; ORGANISM: Human papillomavirus type 16
US-10-530-253-11
```

```
Query Match          100.0%; Score 52; DB 9; Length 248;
```

```

Best Local Similarity 100.0%; Pred. No. 0.058;
Matches 9; Conservative 0; Mismatches 0; Indels 0; Gaps 0;
QY      1 CVYCKQQLL 9
      |||||
Db      127 CVYCKQQLL 135

```

QY	1	CVYCKQQLT	9
Db	127	CVYCKQQLT	135

```

RESULT 9
US-11-192-923A-2
; Sequence 2, Application US/11192923A
; Publication No. US20060018928A1
; GENERAL INFORMATION:
; APPLICANT: PANQ, XIAOMU
; TITLE OF INVENTION: VIRUS-LIKE PARTICLE CONTAINING A DENGUE VIRUS
; TITLE OF INVENTION: RECOMBINANT REPLICON
; FILE REFERENCE: 116620-003
; CURRENT APPLICATION NUMBER: US/11/192, 923A
; CURRENT FILING DATE: 2005-07-29
; PRIOR APPLICATION NUMBER: CN 03115272.4
; PRIOR FILING DATE: 2003-01-30
; PRIOR APPLICATION NUMBER: CN 03115273.2
; PRIOR FILING DATE: 2003-01-30
; NUMBER OF SEQ ID NOS: 45
; SOFTWARE: PatentIn Ver. 3.3
; SEQ ID NO 2
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-11-192-923A-2

```

Query Match	100.0%;	Score 52;	DB 11;	Length 256;
Best Local Similarity	100.0%;	Pred. No. 0.06;		
Matches	9;	Conservative 0;	Mismatches 0;	Indels 0;
				Gaps 0;

QY	1	CVYCKQQL	9
Db	135	CVYCKQQL	143

```

RESULT 10
US-10-530-253-18
; Sequence 18, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530.253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 18
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 35
US-10-530-253-18

```

Query Match	86.5%	Score 45;	DB 9;	Length 149;
Best Local Similarity	87.5%;	Pred. No. 0.63;		
Matches	7;	Conservative	1;	Mismatches 0;
			Indels	0;
			Gaps	0;

QY	1	CVYCKQQL	8
		:	1
Db	30	CVYCKQQL	37

```

RESULT 11
US-10-530-061-1692
; Sequence 1692, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCF/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 1692
; LENGTH: 15
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-1692

```

Query Match	78.8%	Score 41;	DB 9;	Length 15;
Best Local Similarity	75.0%	Pred. No. 0.5;		
Matches	6;	Conservative	2;	Mismatches 0;
			Indels	0;
			Gaps	0;

QY	1	CYCKKQOL	8
		:	1
Db	2	CYCKKEL	9

```

RESULT 12
US-10-530-253-16
; Sequence 16, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McBlhney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530.253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 16
; LENGTH: 149
; TYPE: PRT
; ORGANISM: Human papillomavirus type 31
US-10-530-253-16

```

Query Match	78.8%	Score 41;	DB 9;	Length 149;
Best Local Similarity	87.5%	Pred. No. 3;		
Matches	7;	Conservative	0;	Mismatches 1;
				Indels 0;
				Gaps 0.

QY	1	CVYCKQQL	8
Db	30	CVYCKGQL	37

RESULT 13
US-10-530-253-21
; Sequence 21, Application US/10530253

```
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 21
; LENGTH: 151
; TYPE: PRT
; ORGANISM: Human papillomavirus type 51
US-10-530-253-21
```

```
Query Match      78.8%; Score 41; DB 9; Length 151;
Best Local Similarity 75.0%; Pred. No. 3.1;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
OY      1  CVYCKQQL 8
        |||||:|
Db      30  CVYCKKEL 37
```

```
RESULT 14
US-10-530-253-23
; Sequence 23, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 23
; LENGTH: 155
; TYPE: PRT
; ORGANISM: Human papillomavirus type 56
US-10-530-253-23
```

```
Query Match      78.8%; Score 41; DB 9; Length 155;
Best Local Similarity 75.0%; Pred. No. 3.1;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
OY      1  CVYCKQQL 8
        |||||:|
Db      33  CVYCKKEL 40
```

```
RESULT 15
US-10-530-253-26
; Sequence 26, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
```

```
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 26
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 68
US-10-530-253-26
```

```
Query Match      78.8%; Score 41; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 3.2;
Matches 6; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
OY      1  CVYCKQQL 8
        |||||:|
Db      32  CVYCKRQL 39
```

```
RESULT 16
US-10-530-253-39
; Sequence 39, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 39
; LENGTH: 152
; TYPE: PRT
; ORGANISM: Human papillomavirus
; FEATURE:
; NAME/KEY: MISC FEATURE
; LOCATION: (1)..(152)
; OTHER INFORMATION: where Xaa is any amino acid
US-10-530-253-39
```

```
Query Match      76.9%; Score 40; DB 9; Length 152;
Best Local Similarity 75.0%; Pred. No. 4.6;
Matches 6; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
OY      1  CVYCKQQL 8
        |||||:|
Db      32  CVYCKXZL 39
```

```
RESULT 17
US-11-087-099-2242
; Sequence 2242, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
```

```
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 2242
; LENGTH: 351
; TYPE: PRT
; ORGANISM: Streptococcus agalactiae NEM316
US-11-087-099-2242
```

```
Query Match      73.1%; Score 38; DB 11; Length 351;
Best Local Similarity 75.0%; Pred. No. 20;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 8
          |||||
Db      93 CVYCKHQL 100
```

```
RESULT 18
US-11-087-099-8245
; Sequence 8245, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 8245
; LENGTH: 351
; TYPE: PRT
; ORGANISM: Streptococcus agalactiae 2603V/R
US-11-087-099-8245
```

```
Query Match      73.1%; Score 38; DB 11; Length 351;
Best Local Similarity 75.0%; Pred. No. 20;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 8
          |||||
Db      93 CVYCKHQL 100
```

```
RESULT 19
US-10-453-372-36
; Sequence 36, Application US/10453372
; Publication No. US2006003323A1
; GENERAL INFORMATION:
; APPLICANT: Alsbrook, et al.
; TITLE OF INVENTION: THERAPEUTIC POLYPEPTIDES, NUCLEIC ACIDS ENCODING SAME, AND METHOD
; FILE REFERENCE: 21402-589 A
; CURRENT APPLICATION NUMBER: US/10/453,372
; CURRENT FILING DATE: 2003-06-03
; PRIOR APPLICATION NUMBER: 09/783390
; PRIOR FILING DATE: 2001-02-23
; PRIOR APPLICATION NUMBER: 60/185967
; PRIOR FILING DATE: 2000-03-01
; PRIOR APPLICATION NUMBER: 09/823187
; PRIOR FILING DATE: 2001-03-29
; PRIOR APPLICATION NUMBER: 60/195792
; PRIOR FILING DATE: 2000-03-10
; PRIOR APPLICATION NUMBER: 09/839446
; PRIOR FILING DATE: 2001-03-19
; PRIOR APPLICATION NUMBER: 60/199476
; PRIOR FILING DATE: 2000-03-25
; PRIOR APPLICATION NUMBER: 09/863776
; PRIOR FILING DATE: 2001-05-23
; PRIOR APPLICATION NUMBER: 60/208263
; PRIOR FILING DATE: 2000-05-31
; PRIOR APPLICATION NUMBER: 09/933398
; PRIOR FILING DATE: 2001-08-24
; PRIOR APPLICATION NUMBER: 60/227800
; PRIOR FILING DATE: 2000-08-25
```

```
; Remaining Prior Application data removed - See File Wrapper or PALM.
; NUMBER OF SEQ ID NOS: 1609
; SOFTWARE: CuraSeqlist version 0.1
; SEQ ID NO 36
; LENGTH: 2669
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-453-372-36
```

```
Query Match      71.2%; Score 37; DB 9; Length 2669;
Best Local Similarity 77.8%; Pred. No. 1,4e+02;
Matches 7; Conservative 1; Mismatches 1; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 9
          |||||
Db      2520 CVYCKQOL 2528
```

```
RESULT 20
US-10-530-061-562
; Sequence 562, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SUTHERWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-W
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 562
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-562
```

```
Query Match      69.2%; Score 36; DB 9; Length 10;
Best Local Similarity 75.0%; Pred. No. 2,7;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;
```

```
QY      1 CVYCKQOL 8
          |||||
Db      1 CVYCKATL 8
```

```
RESULT 21
US-10-530-253-20
; Sequence 20, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Cassecci, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 20
```


LENGTH: 158
TYPE: PRT
ORGANISM: Human papillomavirus type 45
US-10-530-253-20

Query Match 69.2%; Score 36; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 23;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 8
|||
Db 32 CVYCKATL 39

RESULT 22
US-11-087-099-7737
Sequence 7737, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 7737
LENGTH: 343
TYPE: PRT
ORGANISM: *Listeria innocua*
US-11-087-099-7737

Query Match 69.2%; Score 36; DB 11; Length 343;
Best Local Similarity 57.1%; Pred. No. 42;
Matches 4; Conservative 3; Mismatches 0; Indels 0; Gaps 0;

QY 1 CVYCKQQL 7
|||
Db 95 CIVCKER 101

RESULT 23
US-11-087-099-3798
Sequence 3798, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 3798
LENGTH: 456
TYPE: PRT
ORGANISM: *Escherichia coli*
US-11-087-099-3798

Query Match 69.2%; Score 36; DB 11; Length 456;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 303 CVYCNKRL 311

RESULT 24
US-11-087-099-6259
Sequence 6259, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement

FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 6259
LENGTH: 456
TYPE: PRT
ORGANISM: *Salmonella enterica* subsp. *enterica* serovar Typhi
US-11-087-099-6259

Query Match 69.2%; Score 36; DB 11; Length 456;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 302 CVYCNKRL 310

RESULT 25
US-11-087-099-9642
Sequence 9642, Application US/11087099
Publication No. US20060041961A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: Genes and Uses for Plant Improvement
FILE REFERENCE: 38-21(53450)B EP
CURRENT APPLICATION NUMBER: US/11/087,099
CURRENT FILING DATE: 2005-03-22
NUMBER OF SEQ ID NOS: 12464
SEQ ID NO 9642
LENGTH: 456
TYPE: PRT
ORGANISM: *Escherichia coli* CFT073
US-11-087-099-9642

Query Match 69.2%; Score 36; DB 11; Length 456;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 302 CVYCNKRL 310

RESULT 26
US-11-188-298-14550
Sequence 14550, Application US/11188298
Publication No. US20060075522A1
GENERAL INFORMATION:
APPLICANT: Abad, Mark S. et al.
TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
FILE REFERENCE: 38-21(53452)B
CURRENT APPLICATION NUMBER: US/11/188,298
CURRENT FILING DATE: 2005-07-22
PRIOR APPLICATION NUMBER: 60/592,978
PRIOR FILING DATE: 2004-07-31
NUMBER OF SEQ ID NOS: 22569
SEQ ID NO 14550
LENGTH: 456
TYPE: PRT
ORGANISM: *Escherichia coli*
US-11-188-298-14550

Query Match 69.2%; Score 36; DB 11; Length 456;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKQQL 9
|||
Db 303 CVYCNKRL 311

RESULT 27

US-11-188-298-19958
; Sequence 19958, Application US/11188298
; Publication No. US2006075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 19958
; LENGTH: 456
; TYPE: PRT
; ORGANISM: Escherichia coli CPT073
US-11-188-298-19958

Query Match 69.2%; Score 36; DB 11; Length 456;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
||| | :|
Db 302 CVYCNRM 310

RESULT 28

US-11-087-099-5321
; Sequence 5321, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 5321
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Escherichia coli K12
US-11-087-099-5321

Query Match 69.2%; Score 36; DB 11; Length 457;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
||| | :|
Db 303 CVYCNRM 311

RESULT 29

US-11-087-099-5761
; Sequence 5761, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 5761
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Salmonella typhimurium LT2
US-11-087-099-5761

Query Match 69.2%; Score 36; DB 11; Length 457;
Best Local Similarity 55.6%; Pred. No. 53;

Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
||| | :|
Db 303 CVYCNRM 311

RESULT 30

US-11-087-099-9118
; Sequence 9118, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 9118
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Escherichia coli
US-11-087-099-9118

Query Match 69.2%; Score 36; DB 11; Length 457;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
||| | :|
Db 303 CVYCNRM 311

RESULT 31

US-11-087-099-10468
; Sequence 10468, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 10468
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Shigella flexneri 2a str. 301
US-11-087-099-10468

Query Match 69.2%; Score 36; DB 11; Length 457;
Best Local Similarity 55.6%; Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CVYCKOQL 9
||| | :|
Db 303 CVYCNRM 311

RESULT 32

US-11-087-099-11070
; Sequence 11070, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 11070
; LENGTH: 457
; TYPE: PRT

```
; ORGANISM: Escherichia coli O157:H7 EDL933
US-11-087-099-11070

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CYYCKOQL 9
   |||||
   :|
Db 303 CVCNSRML 311

RESULT 33
US-11-188-298-4860
; Sequence 4860, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 4860
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Escherichia coli K12
US-11-188-298-4860

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CYYCKOQL 9
   |||||
   :|
Db 303 CVCNSRML 311

RESULT 34
US-11-188-298-8450
; Sequence 8450, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 8450
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Escherichia coli
US-11-188-298-8450

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CYYCKOQL 9
   |||||
   :|
Db 303 CVCNSRML 311

RESULT 35
US-11-188-298-10221
; Sequence 10221, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
```

```
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 10221
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Escherichia coli O157:H7 EDL933
US-11-188-298-10221

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CYYCKOQL 9
   |||||
   :|
Db 303 CVCNSRML 311

RESULT 36
US-11-188-298-16296
; Sequence 16296, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 16296
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Salmonella typhimurium LT2
US-11-188-298-16296

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

QY 1 CYYCKOQL 9
   |||||
   :|
Db 303 CVCNSRML 311

RESULT 37
US-11-188-298-20671
; Sequence 20671, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 20671
; LENGTH: 457
; TYPE: PRT
; ORGANISM: Shigella flexneri 2a str. 301
US-11-188-298-20671

Query Match          69.2% Score 36; DB 11; Length 457;
Best Local Similarity 55.6% Pred. No. 53;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;
```

OY 1 CVYCKOQL 9
| | | | : : |
Db 303 CVYCNRM 311

RESULT 38
US-11-087-099-1964
; Sequence 1964, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 1964
; LENGTH: 465
; TYPE: PRT
; ORGANISM: Yersinia pestis CO92
US-11-087-099-1964

Query Match 69.2%; Score 36; DB 11; Length 465;
Best Local Similarity 55.6%; Pred. No. 54;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 CVYCKOQL 9
| | | | : : |
Db 302 CVYCNRM 310

RESULT 39
US-11-188-298-1940
; Sequence 1940, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 1940
; LENGTH: 465
; TYPE: PRT
; ORGANISM: Yersinia pestis CO92
US-11-188-298-1940

Query Match 69.2%; Score 36; DB 11; Length 465;
Best Local Similarity 55.6%; Pred. No. 54;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 CVYCKOQL 9
| | | | : : |
Db 302 CVYCNRM 310

RESULT 40
US-11-087-099-8259
; Sequence 8259, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B EP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 8259
; LENGTH: 466

; TYPE: PRT
; ORGANISM: Yersinia pestis KIM
US-11-087-099-8259

Query Match 69.2%; Score 36; DB 11; Length 466;
Best Local Similarity 55.6%; Pred. No. 54;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 CVYCKOQL 9
| | | | : : |
Db 303 CVYCNRM 311

RESULT 41
US-11-188-298-18632
; Sequence 18632, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 18632
; LENGTH: 466
; TYPE: PRT
; ORGANISM: Yersinia pestis KIM
US-11-188-298-18632

Query Match 69.2%; Score 36; DB 11; Length 466;
Best Local Similarity 55.6%; Pred. No. 54;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

OY 1 CVYCKOQL 9
| | | | : : |
Db 303 CVYCNRM 311

RESULT 42
US-10-530-061-621
; Sequence 621, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EXS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 621
; LENGTH: 9
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-621

Query Match 67.3%; Score 35; DB 9; Length 9;
Best Local Similarity 83.3%; Pred. No. 1; 9e+05;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

OY 1 CVYCKQ 6
| | | | : : |

Db 4 CVYCK 9

RESULT 43

US-10-530-061-493
; Sequence 493, Application US/1053061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530, 061
; PRIOR FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 493
; LENGTH: 11
; TYPE: PRT
; ORGANISM: Human papillomavirus
US-10-530-061-493

Query Match 67.3%; Score 35; DB 9; Length 11;
Best Local Similarity 83.3%; Pred. No. 4.2;
Matches 5; Conservative 1; Mismatches 0; Indels 0; Gaps 0;

Qy 1 CVYCK 6
|||
Db 6 CVYCK 11

RESULT 44

US-11-004-399-2142
; Sequence 2142, Application US/11004399
; Publication No. US20060053516A1
; GENERAL INFORMATION:
; APPLICANT: Chye, Mee Lee
; APPLICANT: Li, Hong Ye
; APPLICANT: Ramalingam, Sathiskumar
; APPLICANT: Poon, Leo Lit Man
; APPLICANT: Peiris, Joseph Sriyal Malik
; TITLE OF INVENTION: Genetically Modified Plants Comprising SARS-Cov Viral Nucleotide
; FILE REFERENCE: 2587/7316/RDK
; CURRENT APPLICATION NUMBER: US/11/004,399
; CURRENT FILING DATE: 2004-12-03
; PRIOR APPLICATION NUMBER: US 60/527,637
; PRIOR FILING DATE: 2003-12-03
; NUMBER OF SEQ ID NOS: 4043
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 2142
; LENGTH: 35
; TYPE: PRT
; ORGANISM: SARS-Cov Virus
US-11-004-399-2142

Query Match 67.3%; Score 35; DB 11; Length 35;
Best Local Similarity 44.4%; Pred. No. 11;
Matches 4; Conservative 4; Mismatches 1; Indels 0; Gaps 0;

Qy 1 CVYCKQQL 9
|||
Db 15 CVYCKKNT 23

RESULT 45

US-11-188-298-13134

; Sequence 13134, Application US/11188298
; Publication No. US20060075522A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: GENES AND USES FOR PLANT IMPROVEMENT
; FILE REFERENCE: 38-21(53452)B
; CURRENT APPLICATION NUMBER: US/11/188,298
; CURRENT FILING DATE: 2005-07-22
; PRIOR APPLICATION NUMBER: 60/592,978
; PRIOR FILING DATE: 2004-07-31
; NUMBER OF SEQ ID NOS: 22569
; SEQ ID NO 13134
; LENGTH: 109
; TYPE: PRT
; ORGANISM: Oryza sativa
US-11-188-298-13134

Query Match 67.3%; Score 35; DB 11; Length 109;
Best Local Similarity 55.6%; Pred. No. 26;
Matches 5; Conservative 2; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKQQL 9
|||
Db 7 CVYCKSMIL 15

RESULT 46

US-10-530-253-15
; Sequence 15, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 15
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 18
US-10-530-253-15

Query Match 67.3%; Score 35; DB 9; Length 158;
Best Local Similarity 75.0%; Pred. No. 34;
Matches 6; Conservative 0; Mismatches 2; Indels 0; Gaps 0;

Qy 1 CVYCKQQL 8
|||
Db 32 CVYCKTVL 39

RESULT 47

US-10-530-253-19
; Sequence 19, Application US/10530253
; Publication No. US20060014926A1
; GENERAL INFORMATION:
; APPLICANT: Casasetti, Maria C.
; APPLICANT: Smith, Larry
; APPLICANT: Jeffrey K. Pullen
; APPLICANT: Susan P. McElhinney
; TITLE OF INVENTION: HUMAN PAPILLOMAVIRUS POLYPEPTIDES AND IMMUNOGENIC COMPOSITIONS
; FILE REFERENCE: 00630/100M137-US2
; CURRENT APPLICATION NUMBER: US/10/530,253

```
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US2003/031726
; PRIOR FILING DATE: 2003-10-02
; PRIOR APPLICATION NUMBER: US 60/415,929
; PRIOR FILING DATE: 2002-10-03
; NUMBER OF SEQ ID NOS: 65
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 19
; LENGTH: 158
; TYPE: PRT
; ORGANISM: Human papillomavirus type 39
US-10-530-253-19
```

```
Query Match 67.3%; Score 35; DB 9; Length 158;
Best Local Similarity 62.5%; Pred. No. 34;
Matches 5; Conservative 2; Mismatches 1; Indels 0; Gaps 0;
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QY 1 CVYCKQL 8
Db 32 CVYCKRPL 39
```

```
RESULT 48
US-11-087-099-5274
; Sequence 5274, Application US/11087099
; Publication No. US20060041961A1
; GENERAL INFORMATION:
; APPLICANT: Abad, Mark S. et al.
; TITLE OF INVENTION: Genes and Uses for Plant Improvement
; FILE REFERENCE: 38-21(53450)B BP
; CURRENT APPLICATION NUMBER: US/11/087,099
; CURRENT FILING DATE: 2005-03-22
; NUMBER OF SEQ ID NOS: 12464
; SEQ ID NO 5274
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Listeria monocytogenes EGD-e
US-11-087-099-5274
```

```
Query Match 67.3%; Score 35; DB 11; Length 343;
Best Local Similarity 66.7%; Pred. No. 63;
Matches 4; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
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QY 1 CVYCKQ 6
Db 95 CVYCKE 100
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RESULT 49
US-11-045-004-2215
; Sequence 2215, Application US/11045004
; Publication No. US20060078901A1
; GENERAL INFORMATION:
; APPLICANT: BUCHRISSER, CARMEN
; APPLICANT: FRANGEUL, LIONEL
; APPLICANT: COUVE, ELISABETH
; APPLICANT: RUSNIQ, CHRISTOPHE
; APPLICANT: FSIHI, HAFIDA
; APPLICANT: DEHOIX, PIERRE
; APPLICANT: DISSURGET, OLIVIER
; APPLICANT: CHETOUANI, FARID
; APPLICANT: NEJARI, HAFED
; APPLICANT: GLASER, PHILIPPE
; APPLICANT: KUNST, FRANCK
; APPLICANT: COSSART, PASCALE
; APPLICANT: DANIELS, JUSTIN
; APPLICANT: GOEBEL, WERNER
; APPLICANT: KREFT, JURGEN
; APPLICANT: KUHN, MICHAEL
; APPLICANT: NG, EVA
; APPLICANT: VAZQUEZ-BOLAND, ANTONIO
; APPLICANT: DOMINGUEZ-BERNAL, GUSTAVO
; APPLICANT: GARRIDO-GARCIA, PATRICIA
```

```
; APPLICANT: TIERREZ-MARTINEZ, ALBERTO
; APPLICANT: AMEND, ALEXANDRA
; APPLICANT: CHAKRABORTY, TRINAD
; APPLICANT: DOMANN, EUGEN
; APPLICANT: HAIN, THORSTEN
; APPLICANT: BERCHE, PATRICK
; APPLICANT: CHARBIT, ALAIN
; APPLICANT: DURANT, LIONEL
; APPLICANT: PEREZ-DIAZ, JOSE-CLAUDIO
; APPLICANT: BAQUERO, FERNANDO
; APPLICANT: GARCIA DEL PORTILLO, FRANCISCO
; APPLICANT: GOMEZ-LOPEZ, NURIA
; APPLICANT: MADUENIO, ENCARNIA
; APPLICANT: PABLOS, BETRIZ DE
; APPLICANT: WEHLAND, JURGEN
; APPLICANT: KARST, UWE
; APPLICANT: ENTIAN, KARL-DIETER
; APPLICANT: HAUF, JORG
; APPLICANT: ROSE, MATTHIAS
; APPLICANT: VOSS, HAMUT
; TITLE OF INVENTION: LISTERIA MONOCYTOGENES GENOME, POLYPEPTIDES AND USES
; FILE REFERENCE: 05394.0018-02
; CURRENT APPLICATION NUMBER: US/11/045,004
; CURRENT FILING DATE: 2005-01-28
; PRIOR APPLICATION NUMBER: 10/637,657
; PRIOR FILING DATE: 2003-08-11
; PRIOR APPLICATION NUMBER: 10/257,023
; PRIOR FILING DATE: 2002-10-08
; PRIOR APPLICATION NUMBER: PCT/FR01/01118
; PRIOR FILING DATE: 2001-04-11
; PRIOR APPLICATION NUMBER: FR 00/04,629
; PRIOR FILING DATE: 2000-04-11
; NUMBER OF SEQ ID NOS: 2854
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 2215
; LENGTH: 343
; TYPE: PRT
; ORGANISM: Listeria monocytogenes
US-11-045-004-2215
```

```
Query Match 67.3%; Score 35; DB 11; Length 343;
Best Local Similarity 66.7%; Pred. No. 63;
Matches 4; Conservative 2; Mismatches 0; Indels 0; Gaps 0;
```

```
QY 1 CVYCKQ 6
Db 95 CVYCKE 100
```

```
RESULT 50
US-10-530-061-560
; Sequence 560, Application US/10530061
; Publication No. US20060079453A1
; GENERAL INFORMATION:
; APPLICANT: SIDNEY, JOHN
; APPLICANT: SOUTHWOOD, SCOTT
; APPLICANT: SETTE, ALESSANDRO
; TITLE OF INVENTION: HLA BINDING PEPTIDES AND THEIR USES
; FILE REFERENCE: 2060.033US02/EKS/M-M
; CURRENT APPLICATION NUMBER: US/10/530,061
; CURRENT FILING DATE: 2005-04-04
; PRIOR APPLICATION NUMBER: PCT/US03/31308
; PRIOR FILING DATE: 2003-10-03
; PRIOR APPLICATION NUMBER: 60/416,207
; PRIOR FILING DATE: 2002-10-03
; PRIOR APPLICATION NUMBER: 60/417,269
; PRIOR FILING DATE: 2002-10-08
; NUMBER OF SEQ ID NOS: 2503
; SOFTWARE: PatentIn version 3.3
; SEQ ID NO 560
; LENGTH: 10
; TYPE: PRT
; ORGANISM: Human papillomavirus
```

US-10-530-061-560

Query Match 65.4%; Score 34; DB 9; Length 10;
Best Local Similarity 100.0%; Pred. No. 5.9;
Matches 5; Conservative 0; Mismatches 0; Indels 0; Gaps 0;

Oy 1 CVYCK 5
|||
Db 6 CVYCK 10

Search completed: May 5, 2006, 08:29:48
Job time : 10.4 secs

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